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<b>(54) Title:</b> RECEPTOR <b>(57) Abstract</b> <p>A method of screening a chemical for subsequent use as a pharmaceutical agent. The method comprises contacting the chemical with a receptor; and determining whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises EGF-like repeats and/or cadherin-like repeats.</p>		

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## RECEPTOR

The present invention relates to a receptor. In particular the present invention relates to the use of a receptor to screen agents to assess their suitability for subsequent use as pharmaceutical agents, such as therapeutic agents and diagnostic agents.

Receptors are structures that bind chemical stimuli specifically and directly or indirectly transduce a message into the intracellular environment (Watson *et al* 1992 Recombinant DNA Second Edition, Chapter 17, published by Scientific American Books). Some receptors, otherwise known as G-protein coupled receptors (GCRs), are coupled to second messenger systems *via* GTP-binding proteins, otherwise known as G-proteins. The G-proteins connect hormone receptors to adenylate cyclase or other signalling enzymes.

In more detail, the GCRs represent the largest receptor protein class in vertebrates. Typically they are seven-pass transmembrane receptors (see Figure 1). In particular, the GCRs have been shown to be involved in the regulation of a large variety of physiological processes, with many of the genes encoding them mutated in genetic disorders and mutants.

GCRs, can be divided into six families on the basis of their amino acid sequence homologies (similarities) which span across the transmembrane containing region. In this regard, the B family, which is the second largest family, contains the receptors for pituitary adenylate cyclase activating polypeptide, vasoactive intestinal polypeptide (VIP), secretin, growth hormone releasing hormone, diuretic hormone, glucagon, glucagon-like peptide, calcitonin and gastric inhibitory polypeptide.

GCRs can even be placed into functional categories. In this regard, several different types of G-protein coupling have been identified.

For example, the GCR will either interact with an ion channel which is itself a seven-pass transmembrane protein or it can interact with adenylate cyclase, phospholipase C or phospholipase A2, all of which signal to secondary messengers.

- 5 The G-protein can either be stimulatory (Gs) or inhibitory (Gi) and can therefore stimulate or inhibit the action of the ion channel or second messenger pathway they are effecting.

10 All the B family GCRs have been shown to couple to adenylate cyclase *via* a stimulatory G-protein.

In this regard, the present invention provides a new receptor obtainable from animals. The present invention also provides a new use of that receptor.

- 15 Thus, according to a first aspect of the present invention there is provided a method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises EGF-like repeats.

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- According to a second aspect of the present invention there is provided a method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the  
25 receptor comprises cadherin-like repeats.

- According to a third aspect of the present invention there is provided a method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining whether the  
30 chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises EGF-like repeats and cadherin-like repeats.

Preferably, the method includes contacting the chemical-receptor complex with a G-protein and determining whether the chemical-receptor complex stimulates the G-protein.

- 5 Preferably the receptor resembles or is a GCR.

In the following commentary, the term "receptor according to the present invention" includes the receptor as defined in the above-mentioned aspects of the present invention.

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According to a fourth aspect of the present invention there is provided a method of affecting neural development comprising stimulating a receptor with a chemical; wherein the receptor is the receptor according to the present invention. This method can be an *in vitro* or an *in vivo* method.

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According to a fifth aspect of the present invention there is provided the use of the receptor according to the present invention to screen chemicals for subsequent use as a pharmaceutical.

- 20 According to a sixth aspect of the present invention there is provided a chemical that has been screened by the method of the present invention.

According to a seventh aspect of the present invention there is provided a receptor capable of interacting with a G-protein and wherein the receptor comprises EGF-like repeats.

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According to an eighth aspect of the present invention there is provided a protein comprising the amino acid sequence represented as SEQ. I.D. No. 1, or a fragment, homologue or variant thereof.

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According to a ninth aspect of the present invention there is provided a protein comprising the amino acid sequence represented as SEQ. I.D. No. 3, or a fragment, homologue or variant thereof.

- 5 According to a tenth aspect of the present invention there is provided a nucleotide sequence comprising the nucleotide sequence represented as SEQ. I.D. No. 2, or a fragment, homologue or variant thereof, and/or wherein the nucleotide sequence is obtainable from one or more of NCIMB No. 40766, NCIMB No. 40767 and NCIMB No. 40768.

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According to an eleventh aspect of the present invention there is provided a nucleotide sequence comprising the nucleotide sequence represented as SEQ. I.D. No. 4, or a fragment, homologue or variant thereof, and/or wherein the nucleotide sequence is obtainable from one or more of NCIMB No. 40766, NCIMB No. 40767 and NCIMB

15 No. 40768.

According to an twelfth aspect of the present invention there is provided a vector capable of expressing the receptor according to the present invention or the protein according to the present invention, or comprising the nucleotide sequence according

20 to the present invention.

According to a thirteenth aspect of the present invention there is provided a construct comprising or capable of expressing any one of the vector according to the present invention, the receptor according to the present invention, the protein according to the

25 present invention, or the nucleotide sequence according to the present invention.

According to a fourteenth aspect of the present invention there is provided a cell, tissue or organ comprising or capable of expressing any one of the construct according to the present invention, the vector according to the present invention, the receptor

30 according to the present invention, the protein according to the present invention, or the nucleotide sequence according to the present invention.

According to a fifteenth aspect of the present invention there is provided an organism comprising or capable of expressing any one of the cell, tissue or organ according to the present invention, the construct according to the present invention, the vector according to the present invention, the receptor according to the present invention, the protein according to the present invention, or the nucleotide sequence according to the present invention.

According to a sixteenth aspect of the present invention there is provided a receptor capable of interacting with a G-protein and wherein the receptor comprises EGF-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.

According to a seventeenth aspect of the present invention there is provided a receptor capable of interacting with a G-protein and wherein the receptor comprises cadherin-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.

According to an eighteenth aspect of the present invention there is provided a receptor capable of interacting with a G-protein and wherein the receptor comprises EGF-like repeats and cadherin-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.

According to a nineteenth aspect of the present invention there is provided an assay kit comprising a surface having attached thereto or contained within or on any one of the organism according to the present invention, the cell, tissue or organ according to the present invention, the construct according to the present invention, the vector according to the present invention, the receptor according to the present invention, the protein according to the present invention, or the nucleotide sequence according to the present invention.

Typically the assay kit will comprise a series of titre wells capable of holding a suitable sample of the present invention (e.g. the receptor or the gene coding for same

in cells or in a cell free environment). Preferably, the assay kit comprises a series of titre wells, wherein at least one of which well holds a suitable sample of the present invention (e.g. the receptor or the gene coding for same in cells or in a cell free environment). Optionally, the assay kit may comprise one or more G-proteins.

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As mentioned above, if the assay kit of the present invention comprises the receptor of the present invention then that assay kit would be useful for screening chemicals that are capable of interacting with the receptor to form a chemical-receptor complex. With that assay kit, the interaction of the chemical-receptor complex with the G-  
10 protein can be observed either directly or indirectly. An example of indirect observation is observing a change (e.g. an increase) in cAMP levels.

Alternatively, if the assay kit of the present invention comprises the nucleotide sequence of the present invention then that assay kit would be useful for screening  
15 chemicals for affecting expression of that sequence.

Other aspects of the present invention include the use of the receptor of the present invention to screen for agents that are capable of any one or more of

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stimulating the receptor to cause neural cells to divide;

stimulating the receptor to cause neural cells to differentiate;

stimulating the receptor to affect cellular physiology; and

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stimulating the receptor for use in the repair of trauma and/or in the treatment of neurodegenerative diseases.



Other aspects of the present invention include the use of the receptor of the present invention for one or more of:

- stimulating adenylate cyclase;
- 5 increasing cAMP levels; and
- promoting neural growth.

These uses can be *in vitro* or *in vivo* uses.

- 10 Other aspects of the present invention include NCIMB No. 40766, NCIMB No. 40767, and NCIMB No. 40768.

Further aspects of the present invention include:

- 15 a pharmaceutical preparation consisting of or comprising the receptor of the present invention, optionally admixed with or contained within a suitable carrier, diluent or excipient.
- 20 a pharmaceutical preparation consisting of or comprising the nucleotide sequence of the present invention, optionally admixed with or contained within a suitable carrier, diluent or excipient.
- 25 a pharmaceutical preparation consisting of or comprising a chemical when screened by the method of the present invention, optionally admixed with or contained within a suitable carrier, diluent or excipient.
- the use of the receptor according to present invention in the manufacture of a medicament to treat neural disorder.
- 30 the use of the nucleotide sequence according to present invention in the manufacture of a medicament to treat neural disorder.

the use of a chemical when screened by the method of the present invention in the manufacture of a medicament to treat neural disorder.

- 5 A further aspect of the present invention includes a method of treating a subject in need of, or likely to be in need of, treatment for neural disorder wherein the method comprises administering to the subject a receptor according to the present invention, or a protein expressed by the nucleotide sequence according to the present invention, or a chemical when screened by the method of the present invention.
- 10 An additional aspect of the present invention includes a hybrid receptor, and genes coding for the same and vectors etc. comprising same, wherein the hybrid receptor comprises at least a part of the receptor of the present invention and at least a part of another receptor, such as another receptor or even part or all of a G-protein. The hybrid receptor is advantageous as it allows one to affect and/or to tailor the
- 15 stimulation of the receptor to one or more stimuli.

Preferably the receptor is obtainable from neural tissue.

Preferably the receptor is obtainable from ectodermal cells.

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Preferably the receptor is prepared by use of recombinant DNA techniques.

Preferably the receptor is obtainable from deposit NCIMB No. 40766 and/or deposit NCIMB No. 40767.

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Alternatively, the receptor is obtainable from deposit NCIMB No. 40768.

Preferably the receptor comprises the amino acid sequence represented as SEQ. I.D. No. 1, or is a fragment, homologue or variant thereof.

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Preferably the receptor comprises the amino acid sequence represented as SEQ. I.D. No. 3, or is a fragment, homologue or variant thereof.

Preferably the receptor is expressed by the nucleotide sequence represented as SEQ. I.D. No. 2, or is a fragment, homologue or variant thereof.

Preferably the receptor is expressed by the nucleotide sequence represented as SEQ. I.D. No. 4, or is a fragment, homologue or variant thereof.

10 Preferably the chemical is screened to determine if it is useful for one or more of:

- i. causing neural cells to divide;
- ii. causing neural cells to differentiate;
- iii. affecting cellular physiology;
- 15 iv. repairing trauma;
- v. treating neurodegenerative diseases;
- vi. stimulating adenylate cyclase production;
- vii. increasing cAMP levels;
- viii. promoting neural growth.

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Preferred embodiments of the present invention therefore include:

- i. a receptor capable of interacting with a G-protein and comprising EGF-like repeats and/or cadherin-like repeats, wherein the receptor is obtainable from  
25 neural tissue and/or ectodermal cells;
- ii. a receptor comprising the sequence represented as SEQ. I.D. No. 1, or a fragment, homologue or variant thereof capable of being stimulated by a chemical;

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- iii. a receptor comprising the sequence represented as SEQ. I.D. No. 3, or a fragment, homologue or variant thereof capable of being stimulated by a chemical;
- 5 iv. a receptor encoded by the nucleotide sequence represented as SEQ. I.D. No. 2, or a fragment, homologue or variant thereof and wherein the expressed protein is capable of being stimulated by a chemical;
- v. a receptor encoded by the nucleotide sequence represented as SEQ. I.D. No.  
10 4, or a fragment, homologue or variant thereof and wherein the expressed protein is capable of being stimulated by a chemical;
- vi. a receptor obtainable from deposit NCIMB No. 40766;
- 15 vii. a receptor obtainable from deposit NCIMB No. 40767; and
- viii. a receptor obtainable from deposit NCIMB No. 40768.

20 With this aspect of the present invention, the receptor may comprise a plurality of any combination of the features i. to viii. as listed above.

In a highly preferred embodiment the receptor of the present invention is not expressed by the natural genomic DNA sequence when in its natural environment. Thus, highly preferred embodiments include the receptor when prepared by use of recombinant  
25 DNA techniques, including the expression of cDNA or a synthetic nucleotide sequence.

Preferably the receptor is expressed by a cDNA sequence that is obtainable from any one or more of deposits NCIMB 40766, NCIMB 40767 and NCIMB 40768.

In addition, or alternatively, preferably the receptor is expressed by a cDNA sequence that comprises sequence shown as SEQ. I.D. No. 2, or a sequence that is complementary thereto, or a sequence having substantial homology therewith, or a sequence containing any suitable codon substitutions but wherein the resultant protein is capable of acting as receptor as herein defined.

In addition, or alternatively, more preferably the receptor is expressed by a cDNA sequence that comprises sequence shown as SEQ. I.D. No. 4, or a sequence that is complementary thereto, or a sequence having substantial homology therewith, or a sequence containing any suitable codon substitutions but wherein the resultant protein is capable of acting as receptor as herein defined.

In a highly preferred embodiment the nucleotide sequence coding for the receptor of the present invention is not in its natural environment and under the control of the promoter with which it is naturally associated which is also in its natural environment.

Thus, highly preferred embodiments include the use of recombinant DNA techniques using for example cDNA or a synthetic nucleotide sequence.

In a highly preferred embodiment the nucleotide sequence coding for the receptor of the present invention is not in its natural environment and under the control of the promoter with which it is naturally associated which is also in its natural environment, wherein the receptor comprises the amino acid sequence shown as SEQ.I.D. No. 1, more preferably SEQ.I.D. No. 3, or variant, fragment or homologue thereof, wherein the nucleotide sequence is a cDNA sequence that is obtainable from any one or more of deposits NCIMB 40766, NCIMB 40767 and NCIMB 40768, and wherein the nucleotide sequence comprises the sequence shown as SEQ. I.D. No. 2, more preferably SEQ. I.D. No. 4, or a sequence that is complementary thereto, or a sequence having substantial homology therewith, or a sequence containing any suitable codon substitutions but wherein the nucleotide sequence codes for a protein that is capable of behaving as a receptor as herein defined.

Other embodiments of the present invention include: a transformed host organism (such as a microorganism, such as *E. coli*.) capable of producing the receptor according to the present invention as a consequence of the introduction of a nucleotide sequence as herein described; a method for preparing the receptor according to the present invention comprising expressing a nucleotide sequence according to the present invention contained in the host organism and isolating the expressed receptor; and a vector (such as a transformed *pBLUESCRIPT* plasmid, *pGEX* plasmid or *pCDNA3* plasmid) incorporating the nucleotide sequence according to the present invention. By way of example, the receptor can be expressed in *E. coli*, baculovirus, yeast or mammalian cell expression systems.

All of the above-mentioned aspects of the present invention optionally include the combination of the receptor of the present invention with a G-protein.

The term "chemical" includes any chemical compound, including nucleotide sequences both in sense and antisense orientation, proteins, enzymes etc. The term also includes a ligand, wherein a ligand is a natural substance that naturally binds to the receptor.

The term "pharmaceutical agent" includes chemicals for use as diagnostic and/or therapeutic purposes. The term also includes pharmaceutical agents for human and/or veterinary applications.

The terms "screen" and "screening" include the use of the receptor according to the present invention to screen agents to assess their suitability for subsequent use as pharmaceutical agents, such as therapeutic agents and diagnostic agents.

The term "receptor" is used in its normal sense as typically meaning a protein that spans the membrane of a cell and that can bind, on its extra-cellular side, a chemical (otherwise known as a ligand). Binding of the chemical causes changes to the receptor that result in a chemical (enzymatic) reaction being initiated on the intra-cellular part of the receptor. These changes are the first part of a signalling chain of actions that

result in some change to the cells physiology. In the case of GCRs, binding of the chemical causes the receptor to effect a G-protein with which it is associated on the inner membrane surface. This disturbance results in changes to the enzymatic state of the G-protein, which then interacts with the signalling system.

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The term "G-protein" is used in its normal sense as typically meaning a protein which is associated with a receptor and which is capable of being effected by the receptor. Changes in the receptor (binding of chemicals/ligands) effect the enzymatic state of the G-protein and these changes can effect the interactions of the G-protein with other  
10 proteins which are components of a cascade of signalling events. The outcome of the signalling is dependent on the nature of the cell containing receptor and G-protein. In a preferred embodiment, the receptor interacts with a G-protein.

The term "EGF-like repeats" is used in its normal sense as typically meaning a protein  
15 sequence similar to the following "consensus" sequence.

$$CX_{2-6}CX_{4-6}CX_{5-10}CXCX_{8-22}C$$

wherein C is cysteine and X is any amino acid.

20 Preferably, the receptor of the present invention comprises at least one EGF-like repeat and/or at least one cadherin-like repeat. Typically, the receptor of the present invention has between 1 and/or 36 EGF-like repeats and between 1 to 36 cadherin-like repeats. Preferably, the receptor of the present invention has between 1 and 20 EGF-like repeats and/or between 1 and 20 cadherin-like repeats. Preferably, the receptor  
25 of the present invention has between 3 and 10 EGF-like repeats and/or between 3 and 10 cadherin-like repeats. More preferably, the receptor of the present invention has between 4 and 7 EGF-like repeats and/or between 6 and 10 cadherin-like repeats. More preferably, the receptor of the present invention has at least about 6 EGF-like repeats and/or at least about 10 cadherin-like repeats.

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Preferably, the receptor of the present invention comprises at least one EGF-like repeat and at least one cadherin-like repeat.

Preferably, the receptor of the present invention has between 1 and 36 EGF-like repeats and between 1 to 36 cadherin-like repeats. Preferably, the receptor of the present invention has between 1 and 20 EGF-like repeats and between 1 and 20 cadherin-like repeats. Preferably, the receptor of the present invention has between 3 and 10 EGF-like repeats and between 3 and 10 cadherin-like repeats. More preferably, the receptor of the present invention has between 4 and 7 EGF-like repeats and between 6 and 10 cadherin-like repeats. More preferably, the receptor of the present invention has at least about 6 EGF-like repeats and at least about 10 cadherin-like repeats.

The term "chemical-receptor complex" includes binding of the chemical to the receptor, such as by hydrogen bonding and/or covalent bonding. The chemical-receptor complex may then interact with a G-protein. Determination of the formation of the chemical-receptor complex can be achieved by conventional techniques. However, it is preferred to determine formation of the chemical-receptor complex by observing the effect of the complex on a G-protein, such as by observing an increase in cAMP levels.

The term "obtainable from" includes directly or indirectly obtaining the receptor. Examples of indirectly obtaining the receptor include expressing the receptor cDNA by means of a suitable expression system.

The terms "variant", "homologue" or "fragment" include any substitution of, variation of, modification of, replacement of, deletion of or addition of one or more amino acid(s)/nucleic acid from or to the sequence providing the resultant protein is capable of behaving as a receptor as herein defined.

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The expression "substantial homology", which can be otherwise expressed as "substantial similarity", includes homology with respect to structure and/or nucleotide components, providing the resultant protein is a receptor as herein defined.

- 5 With respect to sequence homology (i.e. similarity), preferably there is at least 50 % homology, preferably at least 60% homology, more preferably at least 75% homology, more preferably at least 80% homology, more preferably at least 85% homology, more preferably at least 90% homology, such as at least 95% homology.
- 10 The term "complementary" means that the present invention also covers recombinant nucleotide sequences that can hybridise to the recombinant nucleotide sequences.

The term "construct" - which is synonymous with terms such as "conjugate", "cassette" and "hybrid" - includes all or part of the nucleotide sequence according to the present  
15 invention directly or indirectly attached to another nucleotide sequence, such as a promoter.

The construct may even contain or express a marker which allows for the selection of the genetic construct in the host into which it has been transferred.

20

The construct of the present invention preferably comprises a promoter.

The term "promoter" is used in the normal sense of the art, e.g. an RNA polymerase binding site in the Jacob-Monod theory of gene expression.

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The term "vector" includes an expression vector and a transformation vector. The term "expression vector" means a construct capable of *in vivo* or *in vitro* expression. The term "transformation vector" means a construct capable of being transferred from one species to another - such as from an *E. Coli* plasmid to another host.

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The terms "cell", "tissue" and "organ" include cell, tissue and organ *per se* and when within an organism.

5 The term "organism" in relation to the present invention includes any organism that could comprise the recombinant nucleotide sequence coding for the receptor according to the present invention and/or products obtained therefrom, and/or wherein the recombinant nucleotide sequence according to the present invention can be expressed when present in the organism.

10 Preferably the organism is a transgenic organism. The term "transgenic organism" in relation to the present invention includes any organism that comprises the recombinant nucleotide sequence coding for the receptor according to the present invention and/or products obtained therefrom, and/or wherein the recombinant nucleotide sequence according to the present invention can be expressed within the organism. Preferably  
15 the recombinant nucleotide sequence is incorporated in the genome of the organism.

The term "protein" includes un-modified and modified proteins such as post-translationally modified proteins and glycosylated proteins.

20 The receptor of the present invention is sometimes referred to as the ME2 protein.

The following samples were deposited in accordance with the Budapest Treaty at the recognised depositary The National Collections of Industrial and Marine Bacteria Limited (NCIMB) at 23 St Machar Drive, Aberdeen, Scotland, AB2 1RY, United  
25 Kingdom, on 18 August 1995:

*E. coli* Xl-1 blue containing mouse cDNA plasmid ME2(22) which was allocated deposit number NCIMB 40766;

*E. coli* Xl-1 blue containing mouse cDNA plasmid ME2(78) which was allocated deposit number NCIMB 40767;

30 *E. coli* DH1 containing human cosmid ME2HC20 which was allocated deposit number NCIMB 40768.

These deposits are discussed later in the Experimental Section (see Deposits).

Thus, highly preferred embodiments of the present invention include any one of the  
aforementioned aspects of the present invention but wherein the receptor or the  
5 nucleotide sequence coding for same is obtainable from any one or more of deposits  
NCIMB 40766, NCIMB 40767 and NCIMB 40768.

The present invention will now be described only by way of examples, in which  
reference shall be made to the following Figures, in which:

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Figure 1 is a pictorial representation of a typical GCR;

Figure 2 is a DNA map of the receptor of the present invention;

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Figure 3 is a pictorial representation of the receptor of the present invention;

Figure 4 is a schematic representation of expression patterns;

Figure 5 is a restriction map;

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Figure 6 presents SEQ ID No. 1;

Figure 7 presents SEQ ID No. 2;

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Figure 8 presents SEQ ID No. 3;

Figure 9 presents SEQ ID No. 4; and

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Figure 10 presents an analysis of an amino acid sequence (sequence range 1  
to 2707).

In some of the following commentary, sections (parts) of the gene coding for the receptor of the present invention are sometimes referred to as ME2(x) wherein "x" represents a particular numbered fraction.

5    ME2 Genetic Mapping

Our initial studies revealed that the receptor of the present invention is coded by a single copy gene. This single copy gene is conserved in organisms as diverged as human, mice and fruit flies. In particular, the single copy gene maps to human  
10    chromosome region 22<sup>qter</sup> and mouse chromosome 15. In both of these genomes the receptor gene is contained in a region associated with gastrulation and neural mutants and disorders.

Expression Behaviour

15

To determine the *in vivo* expression of the receptor of the present invention, both reverse transcriptase polymerase chain reaction (RT-PCR) and wholamount *in situ* hybridisation were carried out on wild-type mouse embryos.

20    RT-PCR analysis showed that the receptor of the present invention is first expressed in the early postimplantation embryo between 4 and 6 days post coitum (dpc), then continues until adulthood.

The embryonic expression of the receptor of the present invention precedes the start  
25    of gastrulation, the event which results in the generation of the three germ layers of the developing embryo. Prior to this, the embryo does not contain neural tissue. Embryonic expression of the gene coding for the receptor of the present invention correlates with cells of ectodermal origin, which go on to form the nervous system.

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For ease of reference, Figure 4 is a schematic representation of the expression patterns of the receptor of the present invention in the developing central nervous system, in particular in the developing spinal cord (Figure 4(A)) and the developing hindbrain (Figure 4(B)). In this regard, interesting features of this dynamic expression pattern include the delineation of segments in the developing hindbrain and neural tube. In the hindbrain novel sub-rhombomeric expression was observed. In the neural tube, initially the transcripts are ubiquitous and then resolve into 5,4 and finally 2 dorsoventrally restricted domains, one in the roof plate and one in the floor plate. Gene expression is highly localised and persists throughout development. with adult transcripts localised to the brain and eye.

#### Expression Discussion

Observing the pattern of expression of the receptor of the present invention indicates that it may be involved in the control of neural development. In this regard, the receptor is expressed almost exclusively in neural tissue (which is discussed in more detail later). In particular, expression precedes the first obvious neural structures in the developing embryo and the pattern of subsequent expression is complex. In later embryos, expression around the ventricle of the brain is significant since this is believed to be the region that contains the neural stem cells.

Further observations revealed detection of the receptor of the present invention peri-ventricularly in adult brains. This particular pattern of synthesis is therefore under complex spatial and temporal controls and is the period in which the nervous system is proliferating most rapidly.

Hence, the expression evidence strongly suggests that the receptor of the present invention might play an important part in the "control machinery" of neural development (see later discussion).

Isolation Of The Receptor Of The Present Invention

In some of the following commentary, sections (parts) of the gene coding for the receptor of the present invention are sometimes referred to as ME2(x) wherein "x" represents a particular numbered fraction. The isolation of the complete the receptor of the present invention coding sequence is shown in Figure 5.

In more detail, in step 1 (Figure 5) a mouse 8.5 dpc whole embryo cDNA was screened using a human cDNA clone (16FB2). 16FB2 was originally isolated from a human fetal brain cDNA library by hybridisation with human cosmid ZnFP16 as described in Hoovers *et al.*, (Genomics 10 254-263).

ME2(2) was then isolated from the initial screening of the mouse cDNA library (Figure 5). Extensive sequence analysis of both ME2(2) and 16FB2 has shown that they have sequence homology in a G-rich region in the 3' untranslated region.

Furthermore, complete nucleotide sequencing of ME2(2) showed that it had no homology to any sequences in the publicly accessible DNA databases.

ME2(2) was then used immediately in whole mount *in situ* expression analyses and produced the striking expression pattern of the receptor of the present invention. The remainder of the gene sequence coding for the receptor of the present invention was then isolated as follows.

In step 2 (Figure 5) ME2(2) was used to re-screen the mouse 8.5dpc cDNA library leading to the isolation of 6 different clones, the largest of which being ME2(22). Sequence analysis and database searches with the 3.2kb ME2(22) sequence identified a large open reading frame whose predicted amino acid sequence had homology to the family B group of G-protein coupled receptors. In this regard, ME2(22) covers the region from the polyA tail to trans-membrane region IV.

In step 3 (Figure 5) the 5' EcoRI to *Pst*I fragment of ME2(19) was used to rescreen which led to the isolation of 3 further cDNA clones.

In step 4 (Figure 5) a primer PLKH20 corresponding to sequence 3527 to 3541 in the sequence shown as SEQ.I.D. No. 2 and derived from ME2(42) was used to isolate the H1 fragment using the RACE method (Frohman, M.A. (1993) Methods in Enzymol. 218 340-56). The DNA sequence of H1 was used to identify the DNA from 3295 to 3312 (again see SEQ. I.D. No. 2) which was used to make a new primer, PLKH26, for RACE analysis which gave rise R12 in step 5 (Figure 5).

In step 6 (Figure 5), R12 was used to screen the cDNA library which gave rise to, amongst other clones, the cDNA clone ME2(78) which extends almost to the 5' end.

The minimal set of cDNA clones that defines all of the receptor of the present invention is ME2(78) and ME2(22) (see Figure 2 and Figure 5).

#### DNA Analysis

The DNA sequence of the receptor of the present invention was established entirely using published methods.

In particular, the sequencing methodology used was the Sanger technique (Sanger *et al.*, 1977 Proc. Natl. Acad. Sci. USA. 12 5463-7). The sequencing kits used were supplied by Pharmacia Biotech and the manufacturer's protocols were followed throughout.

DNA was sequenced either by analysis of cloned molecules using sequencing primers specific for vector sequences and sequencing into the cDNA, or by synthesising specific primers, obtained from conventional commercial synthesis companies, and using these to establish DNA sequence directly from internal parts of the cloned cDNA molecule.

The sequence data obtained is shown in the attached Figures as SEQ. I.D. No. 2 (see Figure 7) and SEQ. I.D. No. 4 (see Figure 9). A map of the DNA sequence is represented in Figure 2.

## 5 Re-isolation Of The Receptor Of The Present Invention

The receptor of the present invention was re-isolated by using PCR techniques.

10 Since the gene coding for the receptor of the present invention is over 4.5 kb it is preferable to isolate a cDNA containing the receptor coding regions by a multi-step process, rather than by a one step process using RT-PCR to isolate the whole cDNA. Hence, by using the complete coding sequence for the receptor of the present invention it is possible to isolate a series of cDNA fragments that can then be ligated.

15 In this regard, primers flanking pairs of unique restriction enzyme sites were used to amplify individual regions and subsequent restriction enzyme digestion and ligation to generate the complete sequence. Details of this approach are given below and in Figure 2.

20 **Fragment 1:** Primer pair(3') PLKH31+(5') PLKH52  
Amplification products are digested with *BspHI*.

**Fragment 2:** Primer pair(3') PLKH47+(5')PLKH59c  
Amplification products are digested with *BspHI*

25

Primer	Sequence (5'-3')	Position
PLKH59C	5' CAG CGG GGA CTA CTG CGA GAC TGA AAT	1-27
PLKH47	5' AGC TTG TCG AAG ATG TCA AC	2675-2694
PLKH52	5' ATC TTA CAG CAT GAG AGC CGC C	2414-2435
30 PLKH31	5' GGT AAT GAC ACA GTC ACT GGC ATG	4856-4879



Fragments 1 and 2 were independently amplified from mouse embryonic or adult brain reverse transcribed cDNA under standard PCR conditions. The amplification products were then subsequently directly restriction endonuclease digested with BspHI to give ragged ends. Fragments 1 and 2 were then ligated to each other and then cloned into a T vector.

#### Construction Of A Complete cDNA Clone

Construction of a complete cDNA clone for the gene for the receptor of the present invention was as follows.

In particular, construction of a complete cDNA clone for the gene for the receptor of the present invention was achieved using the protocol detailed in the previous section.

In more detail, the method used relied on the minimal set of cDNA clones obtained from the mouse 8.5dpc libraries mentioned above (see also Figure 2 and Figure 5).

The minimal set of clones ME2(22) and ME2(78) have *EcoRI* linkers and are cloned into pBluescript plasmid vector. Since there is no *EcoRI* site in the ME2 transcript, construction of a complete cDNA clone will be done by *EcoRI* + *AvrII* digestion of ME2(22), isolation of the 2.6kb fragment and ligation of this to the 4.3kb *EcoRI* plus *AvrII* fragment from ME2(78): this product is cloned into the appropriate *EcoRI* digested vector.

#### Deposits

As mentioned above three deposits have been made in accordance with the Budapest Treaty. In this regard, NCIMB 40766 is an *E. coli* XI-1 blue host with a *pBluescript* SK+ vector containing fragment ME2(22) - i.e. nucleotides 3657 to 6794 (see SEQ.I.D.No. 2).

NCIMB 40767 is an *E. coli* XI-1 blue host with a *pBluescript* SK+ vector containing fragment ME2(78) - i.e. nucleotides 1 to 4813 (see SEQ.I.D. No. 2).

NCIMB 40768 is a recombinant cosmid containing the main part of the human  
5 receptor gene according to the present invention. In this regard, the cosmid vector is *pCos2EMBL* and the host cell *E. coli* DH1. The human DNA derives from a region of human chromosome 22<sup>qter</sup> and contains parts of the human receptor of the present invention gene including parts of the 7TM region but not extending further 5' than this.

10

In order to prepare a full length cDNA clone from NCIMB 40766 and NCIMB 40767, the appropriate cDNA fractions can be excised by use of suitable restriction enzymes, isolated and then ligated. The full length cDNA can then be inserted into any suitable expression system and subsequently expressed by suitable means. The *pBluescript*  
15 SK+ plasmids can be recovered from bacterial cells grown in L-broth containing 100 ( $\mu$ g/ml) Ampicillin using routine methods detailed in Sambrook *et al.*, (1989 Molecular cloning - A laboratory manual. Second edition. Cold Spring harbor laboratory press.). Then the receptor cDNA fragments may be isolated subsequent to their excision with *EcoRI*.

20

Likewise, the DNA from NCIMB 40768 and/or fragments thereof can be excised by use of suitable restriction enzymes, isolated, inserted into any suitable expression system and subsequently expressed by suitable means. The DNA can be recovered by growing the bacteria in L-broth supplemented with 30 ( $\mu$ g/ml) Kanamycin and  
25 recovering the DNA according to routine methods detailed in Sambrook *et al.*, (1989 Molecular cloning - A laboratory manual. Second edition. Cold Spring harbor laboratory press). The human DNA fragments can be resolved by cleavage with almost any 6-base recognition enzyme.

30

### Amino acid analysis

The amino acid sequence data are listed in the attached sequence listings as SEQ. I.D. No. 1 (see Figure 6) and SEQ. I.D. No. 3 (see Figure 8).

5

Amino acid analysis of the receptor of the present invention reveals that it appears to be a large membrane spanning receptor having an unusual structure. This structure is pictorially shown in Figure 3.

- 10 In slightly more detail, the C-terminal region appears to have the structure of a 7-pass transmembrane receptor related to the B family of G-protein coupled receptors (GCRs). Thus it is believed that the receptor of the present invention is a new protein.

- 15 Further amino acid analysis reveals that the receptor of the present invention contains EGF-like repeats. In this regard, towards the N-terminus (extracellular) of the receptor there are a number of EGF-like repeats (see Figure 3). One of these EGF-like repeats is ~300 amino acids away from the C-terminal sequence. The EGF-like repeats are shown in Figure 10 (marked "EGF 1" etc.). Divergent EGF-like repeats are also marked.

20

A large number of molecules containing EGF-like repeats have been identified in both vertebrates and invertebrates. These molecules include, for example, blood clotting factors and proteins that are required for correct embryonic development.

- 25 Examples of proteins that are required for correct embryonic development, which molecules have primarily been characterised in invertebrates, include fibropellin, a cell coat protein of sea urchins, glp-1 and lag-12 proteins required for inductive interactions in the nematode worm *C. elegans*, and a number of *Drosophila* proteins including Crumbs, which is required for establishing epithelial cell polarity.
- 30 Additional examples include Notch, Delta and Serrate proteins, which are required for neurogenesis, and Slit, which is a protein involved in axonal pathfinding. Notch and

its ligands Delta and Serrate are involved in cell-cell signalling that determines adjacent cell fates: though this signal is not directly mitogenic.

To date, there have been reports of some isolated vertebrate proteins that have some  
5 homology (similarity) to the invertebrate proteins. For example, three Notch genes, a single Delta, Jagged and Delta-like have been identified to date.

Other vertebrate proteins that have been isolated are EMR-1 (Baud et al 1995 Genomics 26 334-344) and CD97 (Genebank accession no. X84700). However, there  
10 is no mention of possible utility of such proteins. let alone mention of pattern of expression/synthesis.

Further amino acid analysis reveals that the receptor of the present invention contains cadherin-like repeats. These cadherin-like repeats are shown in Figure 10 (marked as  
15 "CD 1" etc.). Cadherin-like repeats have been implicated in protein-protein interactions (Geiger and Ayalon 1992 Ann Rev Cell Biol 8 307-332).

Some of the transmembrane portions of the receptor of the present invention are shown in Figure 10 (marked as "TM 1" etc.).  
20

When the amino acid sequence of the receptor of the present invention is compared with the amino acid sequences of known proteins that are required for correct embryonic development it is observed that there is some sequence homology, though this is less than 80%. More importantly, however, in distinction to the receptor of the  
25 present invention those proteins are all single-pass transmembrane proteins with the cluster of EGF-like repeats in their extracellular domains. In contrast, the receptor of the present invention has a seven-pass transmembrane topology, similar to a GCR.

Accordingly, as there have been no reports in the literature for a receptor found in  
30 neural tissue that is capable of interacting with a G-protein but, in addition, having EGF-like repeats so the receptor of the present invention is novel.

### Functions Of The Receptor Of The Present Invention

The EGF-like repeats and the 7 pass transmembrane (7TM) structures of the receptor of the present invention suggest important functions and utilities for the receptor. In this regard, the EGF-like repeats will bind a ligand which may, as in the case of Notch/Delta, be a protein attached to another cell, or it may be free, as in the case of blood clotting factors. In either case, binding of a ligand will cause the cytoplasmic region to signal to the cellular machinery, *via* a G-protein. Since the 7TM region is most similar to the family B GCRs regions, it is likely that it will signal *via* a stimulatory G-protein which stimulates adenylate cyclase and causes cAMP levels to increase as all family B receptors operate in this fashion.

Likewise, the cadherin-like repeats of the receptor of the present invention suggest important functions and utilities for the receptor.

15

Without wishing to be bound by theory, it is believed that the outcome of this signalling, based upon known examples of GCRs, could be due to one or more of the following effects:

20

1. Stimulation of the receptor of the present invention might cause neural cells to divide.

2. Stimulation of the receptor of the present invention might cause neural cells to differentiate.

25

3. Stimulation of the receptor of the present invention might cause changes to cellular physiology.

Uses

Based on the above-mentioned functions of the receptor of the present invention, it is clear that the receptor can be used in a number of useful utilities. Some of those  
5 utilities are now presented.

1. The receptor of the present invention could be a therapeutic target. In this regard, if the ligand or a modified ligand can be defined, then artificial treatment of neural tissue with this ligand could trigger the receptor of the present invention to signal.  
10 This signal would then trigger the normal response of the receptor of the present invention, which would be a way of modulating the growth, function or properties of brain cells that express the receptor of the present invention. This would therefore have an application in the repair of trauma and in treatment of neurodegenerative diseases, such as Parkinson's disease and Alzheimer's disease.
- 15 2. The receptor of the present invention can be used as a test reagent to identify the ligand. This can be done for example in artificial test systems where the receptor of the present invention is expressed from a recombinant expression vector in cells that otherwise do not express the receptor of the present invention. These cells can then  
20 be used to test for binding of the ligand by studying cAMP level changes upon treatment of cells with proteins or other cells.
3. The nucleotide sequence coding for the receptor of the present invention gene could be used to modify the behaviour of cells by transgenesis. Potential areas of  
25 application include the modification of cells used for transplantation treatments of degenerative diseases and the modification of whole animals by normal transgenic procedures. Both of these applications would result in cells with modified growth potential.

### Screening Protocol

Two screening protocols are now presented.

- 5 The first is based on Lutz *et al.* (1993. FEBS Letters 334, 3-8). In outline, a ME2 expression vector (for example pcDNA-1) is constructed and then introduced into Cos-7 cells by transfection. This results in the cells expressing ME2 and, because of the biological properties of G-proteins, a G-protein becomes naturally associated with ME2 on the cell surface. The cells are then treated *in vitro* with proteins, chemicals,  
10 other cells (intact or broken up). Then one assays for changes to cAMP levels which are caused by ME2 binding a ligand and signalling via the G-protein to alter cAMP levels. If changes are seen, this implies that the ligand is or is contained in, in the substance that was treated with cells. This is the assay of choice for purification of the ligand.

15

- The second method is based upon Cheng & Flanagan (1994, Cell 79 157-168). In this regard, one synthesises in *E. coli* and isolates the N-terminal fragment of ME2 (N terminus of ME2 to the membrane entry point in fig 3) which has been fused at this point to the enzyme alkaline phosphatase. This hybrid protein binds to its normal  
20 ligand. Binding is then detected by looking for the alkaline phosphatase dragged along at its end. This could be used to isolate cDNA clones containing the normal ME2 ligand using exactly the methods detailed in Cheng & Flanagan (1994, Cell 79 157-168).

- 25 The present invention therefore relates to a novel receptor and a novel use of that receptor.

- The nature of the receptor of the present invention, its spatiotemporally restricted expression, coupled with the evolutionary conservation of the receptor gene suggests  
30 that the receptor of the present invention plays a role in the determination of ectodermal cell types within the developing embryo. This has important consequences

as it enables possible pharmaceutical agents to be screened to see if they stimulate the receptor and if so then those agents could be used to promote neural growth. In addition, the receptor can be inserted (such as by way of transplantation or by way of transgenesis of the coding gene) into a subject either in need of treatment or to  
5 develop *in vivo* screening techniques.

Other modifications of the present invention will be apparent to those skilled in the art without departing from the scope of the invention.



CLAIMS

1. A method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining  
5 whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises EGF-like repeats.
2. A method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining  
10 whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises cadherin-like repeats.
3. A method of screening a chemical for subsequent use as a pharmaceutical agent, the method comprising contacting the chemical with a receptor; and determining  
15 whether the chemical interacts with the receptor to form a chemical-receptor complex; wherein the receptor comprises EGF-like repeats and cadherin-like repeats.
4. A method according to any one of claims 1 to 3 wherein the receptor is obtainable from neural tissue.  
20
5. A method according to any one of claims 1 to 4 wherein the receptor is obtainable from ectodermal cells.
6. A method according to any one of the preceding claims wherein the receptor  
25 is prepared by use of recombinant DNA techniques.
7. A method according to any one of the preceding claims wherein the receptor is obtainable from deposit NCIMB No. 40766 and/or deposit NCIMB No. 40767.
- 30 8. A method according to any one of claims 1 to 6 wherein the receptor is obtainable from deposit NCIMB No. 40768.

9. A method according to any one of the preceding claims wherein the receptor comprises the amino acid sequence represented as SEQ. I.D. No. 1, or is a fragment, homologue or variant thereof.
- 5 10. A method according to any one of the preceding claims wherein the receptor comprises the amino acid sequence represented as SEQ. I.D. No. 3, or is a fragment, homologue or variant thereof.
11. A method according to any one of the preceding claims wherein the receptor  
10 is expressed by the nucleotide sequence represented as SEQ. I.D. No. 2, or is a fragment, homologue or variant thereof.
12. A method according to any one of the preceding claims wherein the receptor  
15 is expressed by the nucleotide sequence represented as SEQ. I.D. No. 4, or is a fragment, homologue or variant thereof.
13. A method according to any one of the preceding claims wherein the method  
includes contacting the chemical-receptor complex with a G-protein and determining  
whether the chemical-receptor complex stimulates the G-protein.
- 20 14. A method according to any one of claims 1 to 13 wherein the chemical is screened to determine if it is useful for one or more of:
- i. causing neural cells to divide;
  - 25 ii. causing neural cells to differentiate;
  - iii. affecting cellular physiology;
  - iv. repairing trauma;
  - v. treating neurodegenerative diseases;
  - vi. stimulating adenylate cyclase production;
  - 30 vii. increasing cAMP levels;
  - viii. promoting neural growth.

15. A method of affecting neural development comprising stimulating a receptor with a chemical; wherein the receptor is the receptor as defined in any one of claims 1 to 14.
- 5 16. Use of a receptor as defined in any one of claims 1 to 14 to screen for agents that are capable of stimulating the receptor to cause neural cells to divide.
17. Use of a receptor as defined in any one of claims 1 to 14 to screen for agents that are capable of stimulating the receptor to cause neural cells to differentiate.
- 10 18. Use of a receptor as defined in any one of claims 1 to 14 to screen for agents that are capable of stimulating the receptor to affect cellular physiology.
19. Use of a receptor as defined in any one of claims 1 to 14 to screen for agents that are capable of stimulating the receptor for use in the repair of trauma and/or in the treatment of neurodegenerative diseases.
- 15 20. Use of a receptor as defined in any one of claims 1 to 14 to stimulate adenylate cyclase.
- 20 21. Use of a receptor as defined in any one of claims 1 to 14 to increase cAMP levels.
22. Use of a receptor as defined in any one of claims 1 to 14 to promote neural growth.
- 25 23. Use of a receptor as defined in any one of claims 1 to 14 to screen chemicals for subsequent use as a pharmaceutical.
- 30 24. A chemical when screened by the method according to any one of claims 1 to 15.

25. A receptor capable of interacting with a G-protein and wherein the receptor comprises EGF-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.
- 5 26. A receptor capable of interacting with a G-protein and wherein the receptor comprises cadherin-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.
27. A receptor capable of interacting with a G-protein and wherein the receptor  
10 comprises EGF-like repeats and cadherin-like repeats, wherein the receptor is obtainable from neural tissue and/or ectodermal cells.
28. A receptor according to any one of claims 25 to 27 wherein the receptor is the receptor as defined in any one of claims 4 to 14.
- 15 29. A protein comprising the amino acid sequence represented as SEQ. I.D. No. 1, or a fragment, homologue or variant thereof.
30. A protein comprising the amino acid sequence represented as SEQ. I.D. No.  
20 3, or a fragment, homologue or variant thereof.
31. A nucleotide sequence comprising the nucleotide sequence represented as SEQ. I.D. No. 2, or a fragment, homologue or variant thereof, and/or wherein the nucleotide sequence is obtainable from one or more of NCIMB No. 40766, NCIMB No. 40767  
25 and NCIMB No. 40768.
32. A nucleotide sequence comprising the nucleotide sequence represented as SEQ. I.D. No. 4, or a fragment, homologue or variant thereof, and/or wherein the nucleotide sequence is obtainable from one or more of NCIMB No. 40766, NCIMB No. 40767  
30 and NCIMB No. 40768.

33. A vector capable of expressing the receptor as defined in any one of claims 1 to 14 or the protein of claim 29 or 30, or comprising the nucleotide sequence as defined in claim 31 or 32.
- 5 34. A construct comprising or capable of expressing any one of the vector of claim 33, the receptor as defined in any one of claims 1 to 14 or the protein of claim 29 or 30, or comprising the nucleotide sequence as defined in claim 31 or 32.
- 10 35. A cell, tissue or organ comprising or capable of expressing any one of the construct of claim 34, the vector of claim 33, the receptor as defined in any one of claims 1 to 14 or the protein of claim 29 or 30, or comprising the nucleotide sequence as defined in claim 31 or 32.
- 15 36. An organism comprising or capable of expressing any one of the cell, tissue or organ of claim 35, the construct of claim 34, the vector of claim 33, the receptor as defined in any one of claims 1 to 14 or the protein of claim 29 or 30, or comprising the nucleotide sequence as defined in claim 31 or 32.
- 20 37. An assay kit comprising a surface having attached thereto or contained within or on any one of an organism according to claim 36, the cell, tissue or organ of claim 35, the construct of claim 34, the vector of claim 33, the receptor as defined in any one of claims 1 to 14 or the protein of claim 29 or 30, or comprising the nucleotide sequence as defined in claim 31 or 32.
- 25 38. NCIMB No. 40766.
39. NCIMB No. 40767.
40. NCIMB No. 40768.

41. A method substantially as described herein.
42. A use substantially as described herein.
- 5 43. A receptor substantially as described herein.
44. An amino acid sequence substantially as described herein.
45. A nucleotide sequence substantially as described herein.

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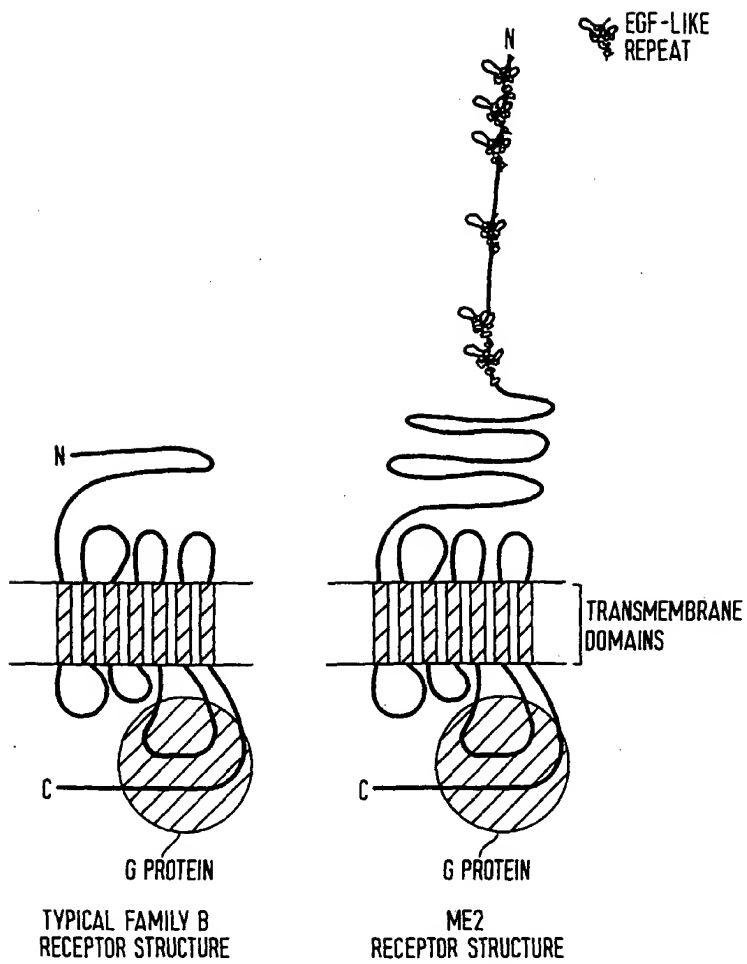
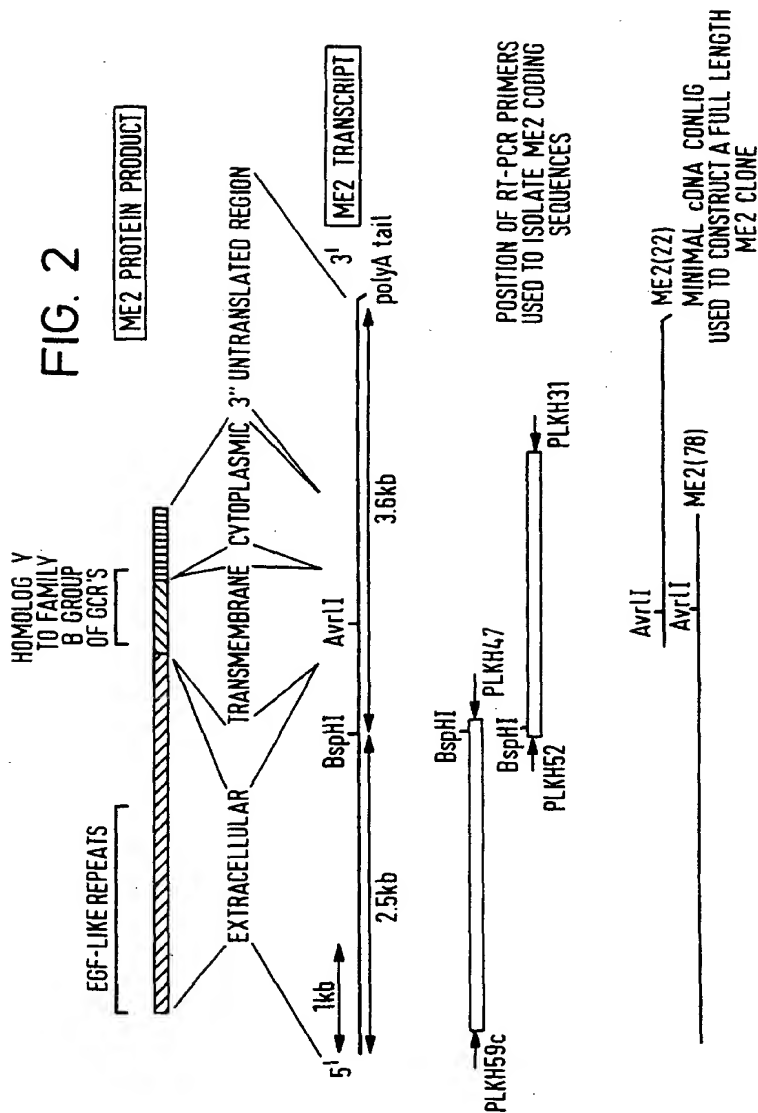


FIG. 1

FIG. 3

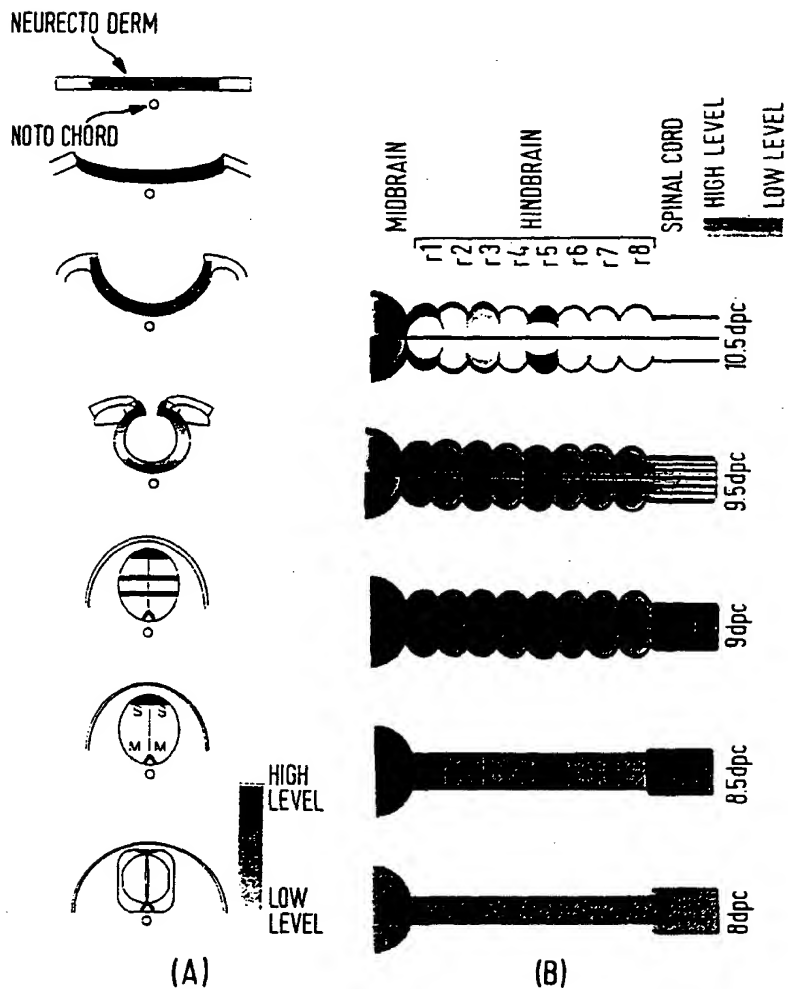
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## FIG. 4

EXPRESSION OF ME2 IN THE DEVELOPING MOUSE  
SPINAL CORD (A) AND HINDBRAIN (B)

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## ME2 TRANSCRIPT WITH POSITIONS OF ALL cDNA CLONES

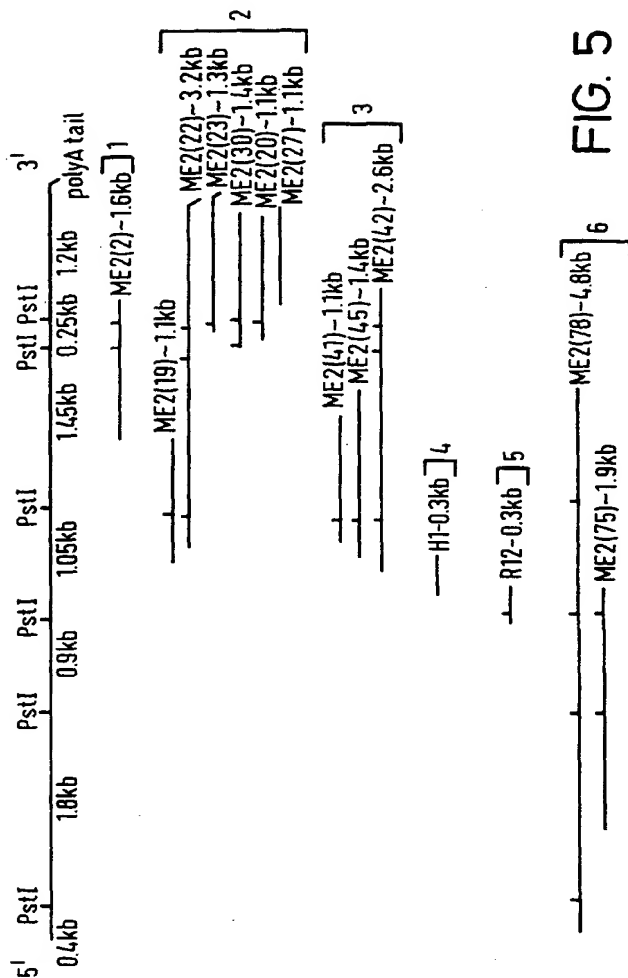


FIG. 5

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AMINO ACID SEQUENCE OF ME2 REGION COVERED BY ME2(78) AND ME2(22)

1573 RESIDUES

GDYCETEIDLCYSNPCGANGGCRSREGGYTCECFEDFTGEHCQVNVRSGRCASGVCK  
NGGTCVNLLIGGFHCVCPPGEYEHYPYCEVSTRSFPPQSFVTFRGLRQRFHFTVSLAF  
ATQDRNALLYNGRFNEKHDFIALEIVEEQQLQTF SAGETTTVTPOVPGGVS DGRW  
HSVLVQYYNKPNIHGLGLPHGPSGEKVAVVTVDCCDAAVAVHFGSYVGNYS CAAGQT  
QSGSKSLDLTGPLLLGGVPNLPEDFPVHSRQFVGCMRNLSIDGRIVDMAAFIANNG  
TRAGCASQRNFCGDTSCONGGTCVNRWNTYLCECLPRFGGKNCEQAMPHPQRFTGES  
VVLWSOLDITISVPWYLGLMFRTRKEDGVLMEATAGTSSRLHLQILNSYIRFEVSYG  
PSDVASMQLSKSRITDGGWHLLIELRSAKEGKDIKYLAVMTLDYGMQDSTVQIGNQ  
LPGLKMRTIVIGGVTEDKYSVRHGFRCMCGVRMGESSTNIATLNMNDALKVRVKDG  
CDVEDPCASSPCPHRPRCDTWDSYSCIDRGYLEKKCVDACLLNPKHVGSLCALP  
NTPRGYSCECGPGHYGQYCEKNVDLPCPKGMMGNRCVAPVTVLSAKALIPTATRPMA  
SARRITTSPOPRIVAFPVTVSPRSRACMDTGQACKPGVIGROCNRCONPFAEV  
TSLGCEVIYNGCPRAFEAGIWWPMKFGQPAAVLCPKGSVGNVVRHCSGEGKWLPE  
L FNCTSGSFVDLKALNEKLNRETRMDGNRSLRLAKALRNATQGNSTLFGNDVRTAY  
QLLARILQHE SRQGFDLAATREANFHEDVVHTGSALLAPATEASWEQIQRSEAGAA  
QLLRHFEAYFSNVARNVKRTYLRPFVITANMILAVDIFDKLNTGAQVPRFEDIQE  
ELPRELESSVSFPADTFKPEKKEGPVVRLTNRRTTPTAQPEPRAERETSSRRRR  
HPDEPGQFAVALVVIYRTLQOLLPEHYDPDRSLRLPNRPVINTPVVSAMVYSEGTP  
LPSSLORPILVEFSLLETEERSKPVCFWNHSLDTGGTGGWSAKGCELLSRNRTHVT  
CQCHSASC AVLMDISRREHGEVLPLKIITYAALSLSLVALLVAFVLLSLVRLTSLN  
LHSTIQEPITHALFFSOLIFMVGINQ TENPFLCTVVAILLHYVSMGTFAWTLVENLHV  
YRMLTEVRNIDTGPMAFYHVVGWGI PAIVTGLAVGLDPQGYGNPDFCWLSLQDTLIW  
SFAGPVGTVIIINTVIFVL SAKVSCQRKHYYERKGVVSMRLTAFLLLLVTATWLL  
GLLAVNSDTLSFHYLFAAFSCLOGIFVLLFYCVANREVRKHLRAVLAGKKLQLDDSA  
TTRATLLTRSLNCNNTYSEGSRHAPHRPGOSTASLDSTTRDEGVOKLSVSSGPARGN  
HGEPDASFIPRNSKKAHGPDOSDSELSLDEHSSSYASSHTSDSEDDGGEAEDKWNP  
AGGPAHSTPKADALANHVPAWNPDES LAGSDSEELDTEPHLKVRPRSAWYSYTGRRRA  
ITVATGPLTRKVGSWPSQWPCLAASPRSSGKAS\*

SEQ ID NO. 1

FIGURE 6

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COMPLETE NUCLEIC ACID SEQUENCE OF ME2 REGION COVERED BY ME2(78) AND ME2(22) cDNA CLONES

NUCLEOTIDES 1 - 6791

CCGGGGACTACTGGGAGACTGAAATTGATCTTTGCTACTCCAATCCGTGCGGGGCCA  
 ATGCGGGCTGCGGAGCCGTGAGGGTGGCTACACTGTGAGTGCTTCGAGGACTTCA  
 CTGGGGAGCATTGCCAGSTGAACGTTCCGCTCAGGCCGCTGTGCCAGGGAGTATGCA  
 AAAACGGGGGCACCTGGCGTGAACCTGCTCATTGGAGGCTTCACTGTGTGTGCCCGC  
 CCGGCGAGTATGAGCATCCCTACTGTGAAGTGAGCACCAGGAGCTTCCACCCCACT  
 CCTTCGTTACCTTCGAGGCCCTGGGCAACGCTTCCACTTACCGCTCCCTGGCGT  
 TTGCCACCAGGACAGGAATGCCCTGCTGCTTACAATGGCCGCTTCAATGAGAAGC  
 ACGACTTCATCGCCCTGSAGATTGTGSAGGAGCAGCTGCAGCTCAGGTTCTCGGCAG  
 GTGAGACCACAACACGGTGACCCGACAGGTTCTGGAGGTGTGAGCGATGGGCGGT  
 GGCATTCCGTGCTGGTGCACTACTACAACAGCCCAACATTGGCCACCTGGGCGCTGC  
 CCCACGGGCGCTGTGAGAGAAAGTGGCTGTGGTGACTGTGGATGACTGTGACGAG  
 CGGTGGCGGTGCACTTTGGAAGTTACGTGGGGAACACAGCTGCGCTGCCAGGGCA  
 CTCAGAGCGGCTCCAAGAAGTCACTGGATCTGACTGGTCCCTCTGCTTCTGGGTGGT  
 TCCCAACCTGCCAGAAGACTTCCCGTGCACAGCCGTCAGTTTGTGGGATGCATGC  
 GAAACCTGTCATCGATGGCGGATTGTGGACATGGCTGCGTTTATTGCCAACATAG  
 GTACAGGGCAGGCTGTGCTTCTCAGAGGAACCTCTCGGATGGGACCTCATGCCAGA  
 ACGGGGGCACCTGTGTGAACAGGTGGAAACAGCTATTATGTAGTGCCCGCTCCGCT  
 TTGTTGGAAAGAACTGTGAACAAGCTATGCCACACCTCAGCGCTTCACTGGTGAGA  
 GCGTCTGTGTGTGGAGTGACCTTGACATCACCATCTCTGTGCTTGGTACCTGGGGC  
 TCATGTTCCGGACCCGGAAGGAGGATGGTGTGCTGATGGAGGACACAGCTGGCACGT  
 TCTCCAGGCTCCATCTCCAGATTCTCAACAGCTACATCCGCTTGGAGTCTCTTAGC  
 GCCCTCTGACGTGGCATCCATGCAGCTGTCCAAGTCCCGGATAAAGTACGGGGGGT  
 GGCATCACCCTGCTCATAGAAGTGAAGAGTGCCAAAGGAGGCAAGGACATCAAAATCC  
 TGGCAGTCATGACCTTGGACTATGGGATGGACACAGCACAGTGACAGATTGGGAATC  
 AGCTTCTCGGGTTGAAGATGCGGACTATTGTTCATCGGAGGTGTGACCGAGGACAAGG  
 TCTCTGTCCGCCATGGTTTCCGAGGCTGTATGCAGGAGTGAGGATGGGAGAGAGCT  
 CCACCAACATTGCCACCCTGAACATGAATGACGCCCTCAAGGTCAGGGTGAAGGAGG  
 GCTGTGATGTGGAGGACCATGTGCTCAAGCCCTGCCCTCCCATAGACCCTGCC  
 GTGACACATGGGACAGCTACTCCTGCATCTGTGACAGAGGCTACTTGGAAAAAAGT  
 GTGTGATGGGTGTCTCTGAACCCCTGCAAGCACGTTGGCAGCTGTGTGCGCTCC  
 CCAACACTCCTCGAGGCTACTCCTGCGAGTGGGACCCGGCCACTATGGGCACTACT  
 GTGAGAAACAAAGTGACCTTCCGTGCCCAAGGCTGGTGGGGAACCGGTGTGTGG  
 CCCCTGTCACTGTGTGTGTCAGCCAGGCTTTGATCCCGACTGCAACAAGACCAATGG  
 CCAGTGCAAGGAGAATTACTACAAGCCCCAGGCCAGGATCGTTGCCCTTCCCTGTGA  
 CTGTTTCCCCCGCTCCACAGCCGTGCTGCGACATGGACACTGGGAGTGTGCT  
 GCAAGCCTGGTGTATCGGCCGTCACTGCAACCGCTGTGATAATCCTTCCCGGAGG  
 TCACCTCGCTCGGCTGTGAAGTGATCTACAATGGGTGTCCAGAGCATTGAGGCTG  
 GCATCTGGTGGCCACAGATGAATTTGGGACGACAGCGGTGCTATGCCCAAAG  
 GATCCGTGGGTAAAGCAGTCCGGCACTGCAGTGGGAGAAAGGCTGGCTTCCCCAG  
 AGCTCTTCAACTGCACCTCTGGCTCCTTTGTGGACCTCAAGGCTTGAACGAGAAAC  
 TGAACCGCAACGAGACAGAATGGACGGGAACCGGTCCTGCGGCTGGCAAGGCTC  
 TGAGGAACGCCACGAGGGGAACAGCACCTCTTTGGCAATGATGTGCGCACCGGCT  
 ACCAGCTTCTGGCCCGCATCTTACAGCATGAGAGCCGCGCAGAGGCTTGAACCTGG  
 CAGCCACCCGAGAGGCTAATTTTCATGAGGATGTGCTCATACAGGCAGCGCCCTCC  
 TGGCCCCAGCTACAGAGGCATCGTGGGAACAGATCCAGCGAAGCGAGGCTGGTGAG  
 CGCAGCTACTGAGGCAATTTGAGGCATATCTCAGCAACGTGGCAGCAAAATGTGAAGA  
 GGACCTATCTGAGGCCCTTGTGTCATGTCACCGCCCAACATGATTTCTGCASTTGACA  
 TCTTCGACAAGCTGAACCTTCAGGGTGCCAGGTGCCAAGGTTTGAGGACATTGAGG  
 AAGAGCTCCAAGGAGCTGGAGTCTCCGTGCTCTCCAGCTGACACCTTCAAGC

SEQ ID NO 2

FIGURE 7

ACCAGAGAAAAAGAGGCCCGCTGGTGAGGCTGACCAACCGAGGACACCCCA  
 CTACGGCAACACCGAGGCCCGAGGCTGAGAGGAGAACTCTCAGCAGCAGGAGGA  
 GACACCCCGATAGGCTGGACAGCTTTGCTGTGCTCGTGTGTCACTTACCGGACCT  
 TGGGTACGCTGTGCTTCAACATGACCCGACATCGAGCTCCGACCTGCCTAT  
 AGCCGCTGTGATCAACACCCCGTGGTGAGTCTGTGTTGACAGTGAAGGAACCC  
 CACTCCCAGCTCTCTGAGAGGCTATCTCTGGTGGAGTTCTCCTGTGGAGACGG  
 AGGAACGAAGCAAACTGTCTGTGTTATCTGGAACAGCTCCCTCGAGTGGTGGGA  
 CTGGAGGGGTGTGTCAGCAAGGGCTGTGAACCTTCTGCGAAGACCGACCACTGCA  
 TCTGCGATCGACCATTTGGCCAGCTGCGGGTGTCTATGCACTATCCAGACGTG  
 ATGCGGGGAGTCTTCCCTCTGAAGTATCACTATGCGCCGCTGTCTGTCT  
 TGGTGGCCCTCTTGGTGGCTTTGTCTCTCTCGCTCGTTCGGACAGTGGCTTCA  
 ACTCGACAGCATCTCCCAAGAACATATCCAGCTGTGTTCTTCTCCAGCTCATCT  
 TCGGCTCGGCATCAACCGACGTGAAGAACGGTTTCTTGACAGTGGTGCCTCAT  
 TCTCTGACATCGTCTTCACTGGGACACTTCCGCTGGACCTTTGGAGAAAGTTCGAT  
 TCTACCGCATGTGACAGAAAGTCGCAACATCGACATGGGCCCTGTCTTCTACC  
 ACGTGGTGGGCTGGGGCATCTCTGCCATTGTACAGGACTGGCTGTTGGCTTGACC  
 CTACGGGCTATGGAAGACCTGATCTTCTGGTGGCTGTCCCTCAGBATAACCTGATT  
 GGAAGCTTTGCTGGGGCTGTGCGAAGCTTATAATCAACACAGTCATCTTTGTGCT  
 TGTCTGCAAGGTTTCTCTGCAAGAAAGACCATTAATGAAGAAAGGGGGTGT  
 TCTCATGCTGAGGACGGCTTCTCTCTGCTGCTGTCTGATCACTGCCACTGCTGCTG  
 TGGGACTGCTGGCGCTCAACGTGACACTTACCTTACTACCTCTTGTCTGCTCT  
 CTAGCTGCTTGCAGGGCATCTTGTCTCTCTGTCTACTGCTGGGCCAACAGGGAG  
 TCGGAAGCACTTAGGGGGGCTGCGGAGGAAGAGCTGCACTGGATGACTCGG  
 CCACCATCTGGGCCATCTTGCTAACCGCGCTCCCTCAACTCAACACACATCAGCG  
 AAGGTTCCAGACATGCTTCGCACTGCGCTTGGGCATGCCAGCTCTTGACAGAT  
 CCACAGGAGATGAAGGGGTCCAGAACTCAGTGTGCTCTTGGGCCACCGCTGGTA  
 ACATGGAGAACACCATGATCATCTTATCCTTAGAACTCCAAAAGCTCAGGCC  
 CTGACTCTGACTCTGACAGTGAAGTGTCTCGGAGAGCAGCATAGTTCTTACGCT  
 TCTCACACATCTCGGACAGGAGATAGTGGCGGAGAGGCTGAAGACAAATGGAAT  
 CGGCTGGGGGCGCCGCATAGCACACCCAAAAGAGATGCTTGGCCACAGCATGTC  
 GAGCTGGCTGGGCCGAGGAGGCTGGCTGGGAGTACAGATGAGGAGTTGACATCT  
 AGCCCACTCTGAAGGTGAGCAGAGGTTCAGCTGGAGTGTACACCGGAGGCGAGGG  
 CAATCACTGTGTCGAGCCCGGCCCTTGACCCGGAAGTGGGTCCTGGCCAGGCAGT  
 GGCCTGTCTTAGCAGCCAGCGCCAGGAGAGCGGGAAGBACCTTGAAAAGAGAG  
 CACTTACCCCGGCCATTTGCCAGAGGACGACATGAAGTCCGGCTGGGAGAGAGC  
 TGGTGATTGTGAGCAGAGCCGCACATCTCCCGACATCTCCTCTTGGCTTGGCG  
 ATGGTGTCTATGACGACTGACTGTGTCAATTACATGAAGCTCAGGAGGGAGGAC  
 GCGGTGACATCTCAATGGGGTGGCATGAATGTACGCCAGGAGGTGCAGGACCA  
 CGGCTGTGACTCAGAGAAACATGAGCAGACTCAACCCACAGACTGCGCGTCAAG  
 CCGTTCAGACTTGAAGCTTGGCTGGGAGTGTGCTTATGGGACAGCGAGGCATTTG  
 TGTAGGGTCTCGCTGATAGCAGCTGGCTTACGACAGGCTCCAGGACGGGAAGCTT  
 TGTACCTCTATAGGAGCTCAGGGCCAGACTGACAGAAAGTGCCAAAGACAGATG  
 TCTTCAGAGAGGAGCTGGAATCTATTAAAGCTGGCATAGCTCGCTCTTAACTAGT  
 AGACAGAAATCAAACTACTGCTACAAGGCACTATGGCCAGAGCTGCACTTGGTG  
 AATCATGTATCGGGCCCTGACTGTGCTCCGAGGCTCTCTTGTGGTACAAGCCC  
 ATACCACAGCTTAGCGGTGCTCTGCAAGGCGCAACCTGGGTTTAAATTTGCTGT  
 CTA AAAATGTAAATAGATATAAATCTCTCTGAGCTGGGAGAGATGGGAGCTGT  
 GTATGCTTTACACTGCTTTGACTCTGCAAGCATGGAGAGCTGAATATGGCATCTCT  
 ACTCTATTGGCCCAAGGAAGCTGCACAGCTTGTACTTGAATCTGGAATGATCAACT  
 AGCTGGTCCAGTGCCTCAGGAGGAGCATTTGGGCTGTGAAGTTGACACAGCTTCT  
 GCAGCTGCTTCTCGGCCAGGAGGTGTGGGTGTTCCGTGGCTTTTGGAGACACGGCA  
 CATGTGCTGATATGTGTGCTCAGCTCATGGATCAGGTCAGGGGCTCAGCTTGTGA  
 GGAAGTGGGCGCACTAGCTATGATTAAGAACTGACCCACTGTCACCCGCCAGCGCC  
 CCCCCCCCCCGGCTCTGATGTCAGTGAAGGCACTTGGAGGCTTTCCGATCAGC

FIGURE 7 CONTINUED

CCAGTGGCAGCGGGAGGACTTCTGCTCCTGCTCCATAAGCTCTCAGAGTGCTCAT  
GTTCTCTGATTCTGAGGGAGCCCAGAGGCTCTCCAACGCTGTACGGTCTTCTCTGCC  
AATGTTGTACTTTTTTCAGACTCCAGGGCTCTCTAGGCAGGCAGGTGCTCCACCCCTG  
AAATCTGACCAATGACATCACTTTTGCTTCAAATGACCAATTGTGCAAGAAACAAG  
CCAGAGTGCCCGTTTTCAATGGTTACCACATTCTTTTTTGGAAATCTCACACCAAGGAC  
CTGTGACCAGCCACTGAGAGCCACGGTGACGCCAAGGCCAAGGGATGGGAGCCTGGA  
GTTACTCAGACACGTTTACTTCAGCATGGACTGTTGTCTGAATCAGGTCCCCAAGT  
ACATGGGGTGACAGTCGCTCGGAATGGAGAACCTCAGGCGAGGCGGTCAAAGGCCA  
GGACTTCTCACCAGCTGTGCTCTCAGAAGTGACAGGACTGCTAGTGACTGACTGG  
TGAGATGAAGCCTACAAGACATGGCCTGGGGTCACGCACTGCTCAGAGGCTGACGG  
GAGGCCCAGGTAGCCCCAAGGATGGCAAGGATACAGAGTGACCTAGCACAGGGGAGCT  
TCAGTCCCAGGTGGTACAGCACCGTGACAACTCCGCAACCCACGCCACCTCAGAAGG  
TGAAGTTTTTGTATCGATCACAACCTATTAGCAAAACAACCCCTGTCAGTTTTAACT  
GTTTTTCTGACCTAAGACTTTCCTTGACCGTAAGACATGGAGATTTTAACAGGTGTA  
TTTATTACTGTTCAACACTGGATGGCAACACAGGTGAGATGATCATCTATAATAAA  
TTAAGATTTTGGATTTGTAAAAA

FIGURE 7 CONTINUED

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## ME2 PROTEIN 1-1798 AMINO ACIDS

NSGLMEKLKQIEEQTKKAQQLLEQOTRRALLEQERKRAGTAVIELRAHPDDEGDAGRLSYQMEALFDER  
 SNGYFLIDAATGAVTTARSLDRETKDTHVLKVSVDHGSPPRASAATYLTVTVDNDHSPVFEQSEYRER  
 IRENLEVGVEYLTRITATGDAPSNNMRYRLLEGAGGVFEIDARSGVVRTRAVVDREEAAEYQLLVEAND  
 QGRNPGPLSASATVHIIVVEDENDNYPOFSEKRYVVOVPEDVAVNTAVLRVQATDRDQGNAAIHYSIVSG  
 NLKGQFYLHSLSGSLDVINPLDFAIREYTLRIKAQDGGRPPLINSSGLVSVQVLVDNDNAPIFVSSPFQ  
 AAVLENVPLGHSVLHIQAVDADAGENARLOYRLVDASTIVGGSSVDSENAPASPOFPFQIHSSGWITV  
 CAELDREEVEHYHSGFEAVDHGSPAMSSASVSITVLDVNDNDPMFTQPVYELRLNEDAAVGGSSVLTIRA  
 RDRDANSVITYQLTGGNTRNRFALSSQSGGLITLALPLDYKQERQYVLAVTASDGRSTRSAQVFINVTD  
 ANTHRPFVQSSHYSVSVSEDRPVGTSIATISATDEDTGENARITYVLEDPVPQFRIDPDTGTITYMTELD  
 YEDQAAATLTAIQAQNGITPKOSDTSLSLEILILDANDNAPRFLRDFYOGSVFEDAPPSTSVLQVSATDRDS  
 GPNGRLTYTQGGDGGDGFYIEPTSGVIRTORRLDRENVAVYNLWALAVDRGSPNPLSASVGIQVSVLD  
 INDNPPVFEKDELELFEVENSPPGVSVARIRANDPDEGPNQIIYQIVEGNVPEVFLDILLSGDLRALVE  
 LDFEVRRDYMVLVQATSAPLVSRATVHIRLLDQNDNPPELPDFQILFNYYVTNKNSFPSPGVIGRIPAH  
 PDLSDSLNYTFLQGNELSLLLLDPATGELQLSRDLNNRPLEALMEVSVSDGHSVTALCTLRTVITITDD  
 MLTNSITVRLNMSQEKFLSPLLFLVEGVATVLSSTKDDIFVFNIQNDTVSSNINLVTSFALLPGGTR  
 GRFFPSEDLOEQIYLRNLTLLTTISAQVLPFDNLCREPCENYMKCVSVLRFDSAPFISSTTVLFRPI  
 HPITGLRCRCPPGFTGDYCETEDLCYSNPGGANGGCRSREGGYTCECFEDTGEHCQVNVSRSGCASGV  
 CKNGGTCVNLIGGFHCVCPPGEYHPYCEVSTRSFPPQSFVTRGLRORFHFTVSLAFATQDRNALLLY  
 NGRFNEKHDFIALEIVEEQQLTFSAGETTTTVPQVPGGVSQGRWHSVLVQYNNKPNIGHGLPHGPSG  
 EKAVAVTVDODCAAVAVHFGSYVGNYSAAQGTQSGSKSLDLTGPLLLGGVPNLPEDFPVHSRQFVCGM  
 RNLSDIGRIVDMAAFIANNGTRAGCASQRNFCOGTSCONGGTCVNRWNTYLCECLRFGGKNCCEQAMPHP  
 QRFTGESVVLWSDLDITISVPWYLGMLFRTRKEDGVLMEATAGTSSRLHLQILNSYIRFEVSYSGSDVAS  
 MOLSKSRITDGGWHLLIELRSAKEGDKIKYLAVMTLDDYMDQSTVQIGNQLPGLKMRITVIGGVTEDEK  
 SVRHGFRGCMQGVMMGESSTNIATLNMNDALKVRVKDGDVEDPCASSPCPPHRPCRDWTDSYSCICDRG  
 YLEKKCVDACLLNPKKHVGSICALPNTPRGYSCEGPGHYGQYCEKNDLPCPKGMWGNRCVAPVTLSA  
 KALIPTATRPASARRITTSQPRIVAFVPTVSPRSHSRACMDTGQACKPGVIGROCNRCNPF AEVT  
 SLGCEVIYNGCPRAFEAGIWWPMKFGQPAAVLCPKGSVGNVAVRHCSGEGKWLPPLEFNCTSGSFVDLKA  
 LNEKLNRENTMRDGNRSLRALAKLRNATQGNSTLFGNOVRTAYQLLARILQHSRQOQFLAATREANFH  
 EDVHTGSALLAPATASWEQIQRSEAGAAQLLRHFEAYFSNVARNVKRTYLRPFVITVANNMILAVDIFD  
 KLNFTGAQVPRFEDIOEELPRELESSVSFPADTFKPEKKEGPPVRLNRRTPTLTAQPEPRAERETSSS  
 RRRRHDPDEPGQFAVALVVIYRTLQQLPEHYDPDHRSLRLPNRPVINTPVVSAMVYSEGTPLPSSLQRP  
 LVEFSLLETEERSKPVCVFWNHSLDTGTTGGWSAKGCELLSRNRTHVTCQCSHSASCAVLMDSRREHGE  
 VLPKITYAALSLSLVALLVAFVLLSLVRTLRSNLHSIPQEPHALFFSOLIFMVGINOTENPFLCTVV  
 AILLHYVSMGTFAMTLVENLHVYRMLTEVRNIDTGPMAFYHVVGWGPATVTLGAVGLDPOGYGNPDFCW  
 LSLQDTLWISFAGPVGTVIIINTVIFVLSAKVSQCKKHYYERKGVVSMRLTAFLLLLTATWLLGLLA  
 VNSDTLSFHYLFAAFSCLOGIFVLLFYCVANREVRKHLRAVLGKKLQLODSDATTATLTLTRSLNCNNTY  
 SEGSRHAPHRPGQSTASLDSTTRDEGVOKLSVSSGPARGNHGEPDASFIPRNSKKAHGPDSDDSELSLD  
 EHSSSYASSHTSDSEDDGGEAEKWNPAAGGAHSTPKADALANHVPAGWPDSELAGSDSELDTEPHLKV  
 RPRSAWYTGRRRAITVATGPLTRKVGSWSPQWPCLAASPRSSGKAS\*

SEQ ID NO. 3

FIGURE 8

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ME2 DNA SEQUENCE 1 - 8210

GAATCCGGGCTGATGAGAAAGCTGAAGCAGATTGAGGAGCAGACTAAGAAGGCTCAGC  
 AAGAGCTGGAAGAGCAGACCCGAGGGCCCTAGAACTTGAGCAGGAACGGAAGCGTGGC  
 GGCACTGCGGTATCGAACTGCGCGGCACGACCCAGACGAAGCGATGCAGGACGCTT  
 TAGCTACCAAGTGGAGGCGCTGTTGATGAGCGCTCTAATGGCTACTTCTCATCGATG  
 CCGCCACGGGTGAGTGACACAGCCCGCTCCCTGGACCGGAGACCAAGGACACTCAT  
 GTACTCAAAGTTAGTGCTGTGGACACGGCTCCCGAGGGCGCTCAGCTGCCACCTACCT  
 CACCGTAACGTGTCAGTGACACTAACGACCACAGCCCGAGTCTTTGAGCAGTCTGAGTATC  
 GAGAGCGAAATTCAGAAAACTGGAGGTGGGCTATGAGGTTCTGACCATCCGTGCCACC  
 GAGCGGGATGCCCTTCCAAACGCAACATGCGCTATCGTCTGCTGAGGGCGCAGGTGG  
 TGTCTTTGAGATAGCGCAGATCAGGTGCTGTCGCGCACACGGGCTGTGGTGGACCGTG  
 AGGAGGCGGTGAGTACCACTGCTGCTGGTGGAGGCCAATGACAGGGTCCGAATCCAGGC  
 CCACCTAGTGCTCAGCCACCGCTCCACATAGTGGTAGAAGACGGAATGACAACTACCC  
 CCAGTTCACTGAGAGAGCGCTATGTGGTTCAAGTCCAGAGAGAGTACCGCTCAACACGG  
 CTGTGCTTCGAGTCCAGGCACTGACCGGGACAGGGGCAGAGTACAGCCATACACTAC  
 AGCATCGTTAGTGGCAACCTGAAGGGTCACTTCTACCTGCATTGCTTATGTTGGAGGCT  
 GATGTTATCAACCGCTGGACTTCGAAGCCATCCGGAATACACCTGGCGCATCAAG  
 CCCAAGATGGGGGCGGCTCCTCTCATTAATTTCTCAGGACTGGTCTCGGTGCGAGGTG  
 TTAGATGTGAACGACAAATGCCGCCATCTTTGTGAGCAGCCCCCTTCAGGCTGCGGTGCT  
 AGAGAAATGTCGCTCGGCACTCAGTCTGTCACATCAAGCGGTGGACGAGATGAGCAG  
 GGGAGAACGCCAGGCTGCACTACCGTCTAGTGGACACAGCCTCCACTATGCTGGGGGGC  
 AGCAGTGTGCACTCTGAGAACCTGCTCTGCCCCAGACTTCCCCTTCCAAATCCACAA  
 CAGCTCCGTTGGATTACTGTGTGCGCGGAGCTGACCGTGAGGAGGTGGAACACTATA  
 GCTTTGGAGTAGAAGCAGTGGACCATGGCTCACCAGCCATGAGCTCTCTGCGAGCGTG  
 TCCATCAGCAGTGTGATGATAATGATAACGACCCCATGTTTACGCGAGCTGTGTATGA  
 GCTGCGTCTGAATGAGGATGCGGCTGTGCGGAGCAGCGTGCTGACCTCAGGGCCGAG  
 ACCGTGATGCCAATAGTGTGATCACTACCAGCTGACGGGTGGGACACCCGCAACCGC  
 TTGCGCACTCAGCAGCAGAGCGCGCTGGCTTATCACTTGGCACTGCCCCGCACT  
 CAAGCAGGAACGCGAGTATGTGCTGGCTGTGACCGCGTCCGATGGCACGCGTTCACACA  
 CCGCGCAGGTCTTTATCAACGTTACAGATGCCAACACCCACAGGCGGTTTTCAGAGT  
 TCCCACTACACGGTCAGTGTGAGTGAAGACGCGCCGTTGGGCACCTCCATCGCTACCAT  
 CAGTGCCACGGATGAGGATACGGGTGAGAACGCCCGCATCACTATGTGCTAGAGGATC  
 CCGTACCACAGTTCGCAATTGACCGGACACTGGCACCATTTATACCATGACGGAATCT  
 GACTATGAGGACCAGGCTGCTACACGCTGGCCATCAGGCTCAGGACAAATGGCATTCC  
 TCAGAACTCAGACACTACCTCTCTGAGATCTCTATCTCGACGCAATGACAAACGCGC  
 CCAGGTTCTGCGAGATTTCTACAGGGTTCTGTTTTGAGGATGCCCCCATCTACCC  
 AGTGTCTCCAGGTCTCTGCTACAGACCGTGACTCAGGCGCTAATGGCCGCTCTCTGTGA  
 CACTTTCCAGGGTGGGATGATGGAGATGGAGATTTCTACATTGAGCCACGCTGCGTG  
 TGATCCGTACCCAGCGCCGCTGGACAGAGAGATGTGGCCGTGTACACACTTTGGGCT  
 CTCGCTGTGGATCGGGGACGCCGAATCCCTCAGTGGCTCAGTGGGAATTCAGGTGAG  
 TGTGTTGGACATTAAACGACAACCCCCAGTGTGTTGAGAAGGACGAGCTGGAGCTGTTTG  
 TGAAGAGAACAGCCCTGTGGGTTCACTGGTAGCAAGAATAAGGGCCACGACCCGAGC  
 GAAGSTCCGAATGCTCAGATCATTTATCAGATCGTGGAGGGCAATGTGCCGAGGCTCT  
 CCAGCTGGACCTACTGAGTGGTGACCTCGTGCCCTGGTCAAGTTGGATTTTGAGGTCC  
 GGAGGGACTATATGTTGGTGGTGACGGCCAGCTCTGCTCTCTGGTAAGCCGGGCCACCC  
 GTGCACATCCGTCTCTGGACAGAAATGACAAACCCAGCGAGTTGGCTGACTTCCAGAT  
 CCTTTTCAACAACTATGTCAACATAAATCAACAGCTTCCCAGTGGTGTGATCGGCC  
 GCATCCAGCCACGACCTTGACCTATCTGACAGCTCAATTACACCTTTCTGCAAGGC  
 AACGAGCTGAGCTGCTGCTGCTGGATCCCGCCACAGGAGAGTTGACGCTCAGCGGGA  
 TCTGGACAACAAACCGGCCACTGGAGGGCTCATGGAGGTGCTGTGTGATGATGTTATCC  
 ACAGCGTCAACGCTCTCTGCACTCTGGCGGTGACCATCATACAGATGACATGCTGACC  
 AACAGCATCACTGTCCGCTGGAGAACATGTGCAAGGAGAAGTTCTGTGCCCGCTGCT  
 GTCCCTCTTTGTAGAAGGGGTGGCCACAGTACTGTCCACCACCAAGGATGACATCTGG  
 TCTTCAACATCCAGAACGACACGACGCTCAGCTCCAACATCTGAACGTGACTTTCTCG  
 GCACTGCTCCCGGTGGCACCGGTGGCCGCTTCTCCGTCAGAGGACCTGCAAGAGAG  
 GATCTACCTGAACCGGACACTGCTCACCACCATCTCCGCCACAGGTGTGCTGCCCTTTG  
 ATGACAACATCTGCTGAGGGAGCCCTGCGAGAATACATGAAGTGGCTGCTGCTGCTGCT

SEQ IN NO.4

FIG 9



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AGGTTTGACAGTTCGGCACCCCTTCATTAGTTCACCACGGTGCTCTCCGGCTATCCA  
 TCCATCACGGGCTGCGCTGCGGCTGCCCGCGGGTTTCACCGGGGACTACTCGCAGA  
 CTCGAATGATCTTTGCTACTCCAATCCGTGCGGGGCAATGGCGGCTGCGGGAGCCGT  
 GAGGGTGGCTACACTTGTGAGTGCTTCGAGGACTTCACCTGGGGAGCATTCGCAGGTGAA  
 CGTTTCGCTCAGGCCGCTGTGCCAGCGAGTATGCAAAAACGGGGACACTCGCTGAACC  
 TGCTCATTTGGAGGCTTCCACTGTGTGTGCCGGCCGCGGAGTATGAGCATCCCTACTGT  
 GAAGTGAGCACCAGGAGCTTCCACCCCAAGTCTTCGTTACCTTCGGAGGCTCGGSCA  
 ACCTCTCCACTTCACCGTCTCCCTGGCGTTTGCCACCCAGGACAGGAATGCCCTGTCTGC  
 TCTACAATGGCGGCTTCAATGAGAAGCACGACTTCATCGCCCTGGAGATTGTGGAGGAG  
 CAGCTGCAGCTCACGTTCTCGGCAGGTGAGACTCACACCAGGTGACACCGCAGGTTCC  
 TGGAGGTGTGAGCGATGGGCGGTGGCATTTCGTGCTGGTGCAGTACTACAACAAGCCCA  
 ACATTGGCCACCTGGGGCTGCCCCACGGGCGCTTGGAGAGAAGGTGGCTGTGGTGAAT  
 GTGATGACTGTGACGACGCGGTGGCGTGCACTTTGGAAGTTACGTGGGGAATACAG  
 CTGCGCTGCCAGGGCACTCAGAGCGGCTCCAAGAAGTCACTGGATCTGACTGGTCTCT  
 TGCTTCTGGGTGGTGTCCCCAACCTGCCAGAGACTTCCCGTGACACGCGTCAAGTTT  
 GTGGGATGCATGCGAAACCTGTCCATCGATGGCCGATTGTGGACATGGCTGCGTTTAT  
 TGCCAAACATGGTACCAGGCGAGGTGTGCTTCTCAGAGGAATCTTGGCATGGGACCT  
 CATGCCAGAACGGGGCACCTGTGTGAACAGGTGGAACACGTAATATGTGAGTGGCCG  
 CTCGCTTGGTGGAAAGAACTGTGAACAAGTATGCCACACCTCAGCGCTTCACTGG  
 TGAGAGCGTGTGTGTGGAGTGACCTTGACATCACATCTCTGTGCTTGGTACCTGG  
 GGCTCATGTTCCGGACCCGGAAGGAGTGTGTGCTGATGAAGCCACAGCTGGCACG  
 TCTTCCAGGCTCCATCTCCAGATTCTCAACAGCTACATCGCTTTGAGGTCTCTCAGG  
 CCCCCTGACGTGGCATCCATGAGCTGTCCAAGTCCCGGATAACTGACGGGGGTGGC  
 ATCACCTGCTCATAGAAGTGAAGAGTGCCAGGAGGGCAAGGACATCAATACCTGGCA  
 CTGACTGACCTTGGACTATGGGATGGACAGAGCAGTGCAGATTGGGAATCAGCTTCC  
 TGGGTGAAGATGCGGACTATTGTATCGGAGGTGTGACCGAGGACAAGGTCTCTGTCC  
 GCCATGGTTTCCGAGGCTGTATGACGGAGTGAGGATGGGAGAGGCTCCACCAACAT  
 GCCACCCTGAACATGAATGACGCCCTCAAGGTGAGGTGAAGGACGGCTGTGATGTGA  
 GGACCATGTGCTCAGAGCCCTGCCCTCCCATAGACCTTCCCTGACACATGGGACA  
 GCTACTCTGTCATCTGTGACAGAGGTAATTTGAAAAAAGTGTGGATGCGTGTCTC  
 CTGAACCCCTGCAAGCAGTGTGGCAGCTGTGTGCGCTCCCAACACTCTCTGAGGCTA  
 CTCTGCGAGTGCAGGCCGGCCACTATGGGAGTACTGTGAGAACAAAGTCGACCTTC  
 CGTGCCCCAAAGGCTGGTGGGGGAACCGGTGTGTGGCCCTGTCACTGTGCTGTGAGC  
 AAGGCTTTGATCCCGACTGCAACAGGACCAATGGCCAGTGAAGGAGAATTACTACAAG  
 CCCCAGCCAGGATCGTTGCCCTTCCCTGTGACTGTTTCCCCCGCTCCACAGCCGTG  
 CCTGGACATGGACACTGGGCAAGTGTGCTGCAAGCCTGGTGTCACTGGCCGTGAGTGC  
 AACCCTGTGATAATCCTTTTGGCGAGGTCACTGCTCGGCTGTGAAGTATCTACAA  
 TGGGTGTCCAGAGCATTTGAGGCTGGCATCTGTGGCCACAGATGAATTTGGGCAGC  
 CAGCAGCGGTGCTATGCCCCAAGGATCCGTGGGTAACGAGTCCGCGACTGCAAGTGGG  
 GAGAAAGGCTGGCTTCCCCAGAGCTCTTCAACTGCACCTTGGCTCCTTTGTGGACCT  
 CAAGGCTTTGAACAGAGAACTGAACCGCAACGAGACAAGATGAGCGGGAACCGGTCCC  
 TGGCGCTGGCAAGGCTCTGAGGAACGCCACGAGGGGAACAGCACCTCTTTGGCAAT  
 GATGTGCGCAGCGCTACAGCTTCTGCGCCGCACTCTACAGCATGAGAGCCGCGAGCA  
 GGGCTTTGACCTGGCAGCCACCAGAGGCTAATTTTCATGAGGATGTGCTCCATACAG  
 GCAGGCGCTCTTGGCCCCAGCTACAGAGGATCTGTGGGAACAGATCCAGCGAAGCGAG  
 GCTGGTGACGCGAGCTACTGAGGCAATTCGAGGCACTACTTCAGCAACGTGGCAGGAA  
 TGTGAAGAGGACCTATCTGAGGCCCTTCTGTCATGTCACCGCCAAACATGATTTTCAG  
 TTGACATCTTCACAAGCTGAACCTCACGGGTGCCAGGTGCCAAGGTTTGAAGGACATT  
 CAGGAAGAGCTCCCAAGGGAGCTGGAGTCTCTCGTCTCTTCCAGCTGACACCTTCAA  
 GCCACAGAGAAAAAAGGCCCCGTGGTGAGGTGACCAACCGGAGGACTACCCAC  
 TCACGCAACAACAGAGCCAGGGCTGAGAGGGAAACCTCATCCAGCAGAGCGAGGAGA  
 CACCCGATGAGCTGGAGCACTTTGCTGTGGCTGGTGTGCTATTACCGGACCTGGG  
 TCAGTGTCTGCTGAACACTATGACCCCGACCATCGCAGCTCCGACTGCTTAACCGGG  
 CTGTTCATCAACACCCCGGTGGTGAAGTGTATGGTGTACAGTGAGGGAACCCCTCC  
 AGCTCTCTGAGAGGCTATCTGTTGGAGTCTCTCTGTTGGAGCGGAGGAACGAG  
 CAACCTGTCTGTGATCTTGAACCACTCCCTGACACTGGTGGGACTGGAAGGTGGT  
 GACCCAGAGGCTGTGAACCTCTGTGAGGAACCGGACCCAGCTCACTTGCAGTGAGC  
 CATTCGGCAGCTGCGCGGTGCTCATGGACATTCAGACGTGAGCACGGGAGGTTCT

FIGURE 9 CONTINUED

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GCCCTGAAGATCATCACCATGCCGCCCTGTCTTGTCTTTGGTGGCCCTCCTGGTGG  
 CCTTTGTCTTCTCTCGCTGGTTCCGACACTGCGCTCCAACTGCACAGCATCCACAAA  
 GAAGCTATCCAGCTCTGTTCTTCTCCAGCTCATCTTCATGGTGGCATCAACCAGAC  
 TGAGAACCCGTTTCTCTGCACAGTGGTGGCATCTCTCTGCACTACGTCTCCATGGCA  
 CCTTCGCTGGACCTTTGTGGAGAACTTGCAATGTCTACCGCATGCTGACAGAAAGTGGC  
 AACATCGACACTGGGCCCATTGGCGTTCTACACAGTGGTGGCTGGGGCATCCCTGCCAT  
 TGTACAGGACTGGCTGTTGGCTTGGACCTCAGGGCTATGGAAACCCGTGACTTCTGCT  
 GGCTGTCCCTTCAGGATACCCTGATTTGGAGCTTTGTCTGGGCTGTCTGGAAAGGTTATA  
 ATCATCAACACAGTATCTTTGTCTGTCTGCAAGGTTTCTGCAAGAAAGACCA  
 TTATTATGAAAGAAAGGGGTTGTCTCATGCTGAGGACGGCTTCTCTGCTGCTGCTG  
 TCGTCACTGCCACTGGCTGCTGGGACTGCTGGCGGTCAACAGTGACACTCTTAGCTTT  
 CACTACCTCTTTGCTGGCTTCAGCTGCTTGCAGGGCATCTTTGTCTCTGTTCTACTG  
 CGTGGCAACAGGGAGGTGCGGAAGCACTTGAGGGCGGTGTGGCAGGGAAGAGTGC  
 AGCTGGATGACTCGGCCACCACTCGGCCCACTCTGCTAACGCGTCCCTCACTGCAAC  
 AACACTACAGCGAAGGGTGCAGACATGCTCCGCAACGCGCTTGGGCACTCCACAGCTC  
 TCTGGACAGTACCACAGGATGAAGGGTCCAGAAACTCAGTGTGCTCTTGGGCCAG  
 CCCGTGGTAACCATGGAGAACAGATGCATCTTCACTCTAGGAACTCCAAAAAGCT  
 CACGGCCCTGACTCTGACTCTGACAGTGAAGCTGTCTTGGAGAGCAGATAGTTCCTA  
 CGCTCTTCCACACATCGGACAGCGAGGATGATGGCGGAGGGTGAAGACAAAGTGA  
 ATCCGGCTGGGGCCCCGCCATAGCACCCAAAAGCAGATGCTCTGGCCAAACACGTC  
 CCAGCTGGCTGGCCGACGAGAGCTGCTGGTGGGAGTGACAGTGAGGAGTTGGACACTGA  
 GCCCACTTGAAGGTGAGACCAAGGTGAGCTGGAGTTACACCGGCGAGGGCAAGGCA  
 TCACCTGGCGACCGGCCCTTGACCCGGAAAGTGGGGTCTGGCCAGGCAAGTGGCGG  
 TGCTTAGCAGCCAGCCCCAGGAGCAGCGGAAAGGCATCTGAAAAACAAAGTCACTAC  
 CCGCGGGCATTGCCAGAGCAGCACTGAAGTCCCGCTGCGAGAGAAGCTGGCTGATT  
 GTGAGCAGAGCCCCACATCTCCCGCACATCTCTCTTGGCTCTGGGATGGTGTCCAT  
 GCGACTGACTGTGTCTATTACCATCAAGACTCCGAGGAGGGAGCCAGGCGTGAGCATCT  
 CAATGGGGTGGCATGAATGTAGCCACAGGGAGTGCAGGCCAACGTTCTGACTAGA  
 GAAACCATGAGGACAGCTCAACCCACAGACTGCGGGTCAAGCCCTCAGACCTTGACAGA  
 CTGCTGGGACTGCTGCCATGCGGACAGGAGGCACTTGTGTGAGGGTCCCTGGCCATA  
 GCAGCTGGCTCTACGACAGCCGTCCAGACGGGAAGCCCTTGACCTCATAGSAGCTCAG  
 GGGCCAGACTCTGACAAAGTGCCAAAGCCACAGATGTCTCCAGAGGGAGAGCTGCACTT  
 CATTAAGGCTGGCATAGCTCCGTCTTAGACTGAAGACAGAATCCAAACCATGCTGTAC  
 AAGAGGCCATTGAGCCAGAGCTGGACTTGGTGAATCATTGTACGGGGCCCTTCAACTGT  
 CCGCAGGGCTCTCTTGTGGTACAAGCCATCACCACAGGCTAGCGGTGCTCTGCA  
 ACGGCAACCTGGGTTTTAAATTGCTGTCTAAAAATGTAAATAGATATAAATCTCTCCC  
 TGGACTTGGGAGAAGTGGGAGCTGTGTATGCTTTACACTGCTTTGACTCTGCAGCCAC  
 TGGAGAGCCATGAATGGCATCTCACTCTATTGGCCAAAGGAAGCTTGACAGTTGACT  
 TGAATCTGGAATGAGTCACTCAGCTGGTCCAGTGCCAGTAGGGAGCATATGGGCT  
 GTGAAGTTGACACAGCTTCTGCAGCTGCTTCCGGCAGGGAGTGTGGGTGTTCCGCTGG  
 CCTTTGGAGCACACGGCACATGTGCTGCGTATGTGTGCTCAGCTCCATGGATCCAGGT  
 CAGGGCTCACCTGTGAGGAGTGGGGCCATAGCTATGATATGAAGTCTGACCACCCGT  
 CCACGCCACGGCCCCCCCCCCCCCGCTCTGATGCACTGAGGCGACTCTGGAGCT  
 TTCCAGTCAGCCAGTGCGCAGCGGAGGACTTCTGCTCTGCTCTCCATAAGCTCTCA  
 GAGTGTCTATGTTCTGATTCTGAGGGAGCCAGAGGTCTTCCAAAGCTGTACAGGCTCC  
 TTCTGCCAATGTTGACTTTTTTACAGACTCCAGGGCTCTCTAGGCGAGGAGTGTCTCCA  
 CCCCATAATCTGACCAATGACATCACTTTGCTTCAAATGACCAATTGTGCAAGAGAAAC  
 AAAGCCAGAGTGCCGTTTTCAATGGTTACCACTTTCTTTTGAATCTCACCAAGG  
 ACCTGTGACAGGCACTGAGAGCCACGCTGCAGCCAAAGGCAAGGATGGAGCTGGA  
 GTTACTCAGACAGTTTACTTCAGCATGGACTGTTGTCTGAATCAGGTCCCCAAGTAC  
 ATGGGGTGACAGTGCCTCGAATGGAGAACCTCAGGCGAGGGGTCAAAGGCCAGGAC  
 TTCTACCAAGCTGTGCTCTCAGAAGTGCACAGGACTGCTAGTGACTGACTGGTGAGAT  
 GAAAGCTTCAAGACATGGCTTGGGGTCAAGCACTGCAAGAGGCTGACGGGAGGCCA  
 GGTAGCCCAAGGATGCCAAGGATACAGAGTGACCTAGCACAGGAGGCTTCACTCCAG  
 GTGGTACAGCACCGTGACAACTCCGCAACCCACGCGACCTCAGAAGGTGAAGTTTTGA  
 TCGATTCACTAATTATGCAAAACAAACCTGTCAAGTTTAAACTGTTTTCTGACCTA  
 AGACTTCTTACCGCTAAGACATGGAGATTTTAAACAGGTGATTTATTACTGTTACG

FIGURE 9 CONTINUED

ACTGGATGGCAACACAGGTGAGATGATGCATCTATAATAAATTAAGATTTTGGATTG  
TAAA

FIGURE 9 CONTINUED

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## FIG. 10

	5	10	15	20	25	30	35	40	45	50
	NSGLM	EKLKQ	IEEQT	KKAQQ	ELEEQ	TRRAL	ELEQE	RKRAG	TAVIE	LRAHD
	55	60	65	70	75	80	85	90	95	100
CD1	PDEGD	AGRLS	YQMEA	LFDER	SNGYF	LIDAA	TGAVT	TARSL	DRETK	DTNVL
	105	110	115	120	125	130	135	140	145	150
	KVSAV	DHGSP	RRSAA	TYLTV	TVSDT	NDHSP	VFEQS	EYRER	IRENL	EVGYE
	155	160	165	170	175	180	185	190	195	200
CD2	VLTR	ATDGD	APSNA	NMRYR	LLEGA	GGVFE	IDARS	GVVRT	RAVVD	REEAA
	205	210	215	220	225	230	235	240	245	250
	EYQLL	VEAND	QGRNP	GPLSA	SATVH	IVVED	ENDNY	PQFSE	KRYVV	QVPED
	255	260	265	270	275	280	285	290	295	300
CD3	VAVMT	AVLRV	QATDR	DQGQN	AAIHY	SIVSG	NLKGQ	FYLHS	LSGSL	DVINP
	305	310	315	320	325	330	335	340	345	350
	LOFEA	IREYT	LRIKA	QDGR	PFLIN	SSGLV	SVQVL	DVNDN	APIFV	SSFFQ
	355	360	365	370	375	380	385	390	395	400
	AAVLE	NVPLG	HSVLI	IQAVD	ADAGE	NARLQ	YRLVD	TASTI	VGGSS	VDSEN
CD4	405	410	415	420	425	430	435	440	445	450
	PASAP	DFFPQ	IHNSS	GWITV	CAELD	REEVE	HYSFG	VEAVD	HGSPA	MSSSA
	455	460	465	470	475	480	485	490	495	500
	SVSIT	VLDVN	DNDPM	FTQPV	YELRL	NEDAA	VGSSV	LTLRA	RDRDA	NSVIT
	505	510	515	520	525	530	535	540	545	550
CD5	YQLTG	GNTRN	RFALS	SQSGG	GLITL	ALPLD	YKQER	QYVLA	VTASD	GTRSH
	555	560	565	570	575	580	585	590	595	600
	TAQVF	INVTD	ANTHR	PVFQS	SHYTV	SVSSE	RPVGT	SIATI	SATDE	DTGEN
	605	610	615	620	625	630	635	640	645	650
CD6	ARITY	VLEDP	VPQFR	IDPDT	GITYT	MTELD	YEDQA	AYTLA	ITAQD	NGIPQ
	655	660	665	670	675	680	685	690	695	700
	KSDTT	SLEIL	ILDAN	DNAPR	FLRDF	YQGSV	FEDAP	PSTSV	LQVSA	TDRDS
	705	710	715	720	725	730	735	740	745	750
CD7	GPNGR	LLYTF	QGGDD	GDGDF	YIEPT	SGVIR	TQRRL	DRENV	AVYNL	WALAV
	755	760	765	770	775	780	785	790	795	800
	DRGSP	NPLSA	SVGIQ	VSVLD	INDNP	PVFEX	DELEL	FVEEN	SPVGS	VVARI

# FIG. 10 CONTINUED

805 810 815 820 825 830 835 840 845 850  
C08 RANDP DEGPN AQIYI QIVEG NVFEV FQDL LSGDL RALVE LDFEV RRDYM  
855 860 865 870 875 880 885 890 895 900  
LVVQA TSAPL VSRAT VHRL LDQND NPPEL PDFQI LFNNY VTKNS NSFPS  
905 910 915 920 925 930 935 940 945 950  
C09 GVIGR IPARD PDLSD SLNYT FLOGN ELSLL LLDPA TGELO LSRDL DNNRP  
955 960 965 970 975 980 985 990 995 1000  
LEALM EVSVS DGIHS VTALC TLRVT IITDD MLTNS ITVRL ENMSQ EKFLS  
1005 1010 1015 1020 1025 1030 1035 1040 1045 1050  
PLLSL FVEGV ATVLS TTKDD IFVFN IQNDT DVSSN ILNVT FSALL PGGTR  
1055 1060 1065 1070 1075 1080 1085 1090 1095 1100  
GRFFP SEDLO EQIYL NRILL TTISA QRVL PDDNI CLREP CENYM KCVSV  
1105 1110 1115 1120 1125 1130 1135 1140 1145 1150  
LRFDS SAPFI SSTTV LFRPI HPITG LRCRC PPGFT GDYCE TEIDL CYSNP EGF1  
1155 1160 1165 1170 1175 1180 1185 1190 1195 1200  
CGANG GCRSR EGGYT CEGFE DFTGE HQVNV VRSGR CASGV CKNGG TCNVL EGF2  
1205 1210 1215 1220 1225 1230 1235 1240 1245 1250  
LIGGF HGVCP EGEYE HPYCE VSTRS FPPQS FVTER GLRQR FHFTV SLAFA  
1255 1260 1265 1270 1275 1280 1285 1290 1295 1300  
TQDRN ALLLY NGRFN EKIDF IALEI VEEQL QLTFS AGETT TTVTP QVPGG  
1305 1310 1315 1320 1325 1330 1335 1340 1345 1350  
VSDGR WHSVL VQYYN KPNIG HGLLP HGPFG EKVAV VTVDD CDAAV AVHFG  
1355 1360 1365 1370 1375 1380 1385 1390 1395 1400  
SYVGN YSCAA QGTQS GSKKS LDLTG PLLLG GVPNL PEDFP VHSRO FVGCM  
1405 1410 1415 1420 1425 1430 1435 1440 1445 1450  
RNLSE DGRIV DMAAF IANNG TRAGC ASQRN FCDGT SCQNG GTCVN RWNFY  
1455 1460 1465 1470 1475 1480 1485 1490 1495 1500 EGF3  
LCEP LRFGG KXCEQ AMPHP QRFTG ESVVL WSDLD ITISV PNYLG LMFT  
1505 1510 1515 1520 1525 1530 1535 1540 1545 1550  
RKEDG VLMEA TAGTS SRLHL QILNS YIRFE VSYGP SDVAS MQLSK SRITD  
1555 1560 1565 1570 1575 1580 1585 1590 1595 1600  
GOWH LLIEL RSAKE GKDIK YLAVM TLDYG MDQST VQIGN QLPGL RMRTI  
1605 1610 1615 1620 1625 1630 1635 1640 1645 1650  
VIGGV TEDRV SVRHG FRGCM QGVFM GESST NIATL NMNDA LRVRV KDGCD

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## FIG. 10 CONTINUED

1655 1660 1665 1670 1675 1680 1685 1690 1695 1700  
 EGF4 VEDPC ASSPC PPHRP CRDTW DSYSC EGF4 EGF4 EGF4 EGF4 EGF4  
 1705 1710 1715 1720 1725 1730 1735 1740 1745 1750  
 EGF5 LCALP NTPRG YSCPC GPGHY GOYCE NKVDL PCPRG WNGNR CVAPV TVLSA  
 1755 1760 1765 1770 1775 1780 1785 1790 1795 1800  
 KALIP TATRP MASAR RITTS POPRI VAFPV TVSPR SHSRA CDMOT QQCAC  
 1805 1810 1815 1820 1825 1830 1835 1840 1845 1850  
 KPGVI GROCJ RCDNP FAEVT SLOCE VIYNG CPRAF ENGIW WPQMK FGQPA  
 1855 1860 1865 1870 1875 1880 1885 1890 1895 1900  
 AVLCP KGSVG NAVRH CSGEK GWLPP ELFNC TSGSF VDLKA LNEKL NRNET  
 1905 1910 1915 1920 1925 1930 1935 1940 1945 1950  
 RMDGN RSLRL AKALR NATOG NSTLF GNDVR TAYQL LARIL QHESR QOGFD  
 1955 1960 1965 1970 1975 1980 1985 1990 1995 2000  
 LAATR EAMFH EDVVH TGSAL LAPAT EASWE QIQRS ENGAA QLLRH FEAFX  
 2005 2010 2015 2020 2025 2030 2035 2040 2045 2050  
 SNVAR NVKRT YLRPF VIVTA NMILA VDIFD KLNFT GAQVP RFEDI OEELP  
 2055 2060 2065 2070 2075 2080 2085 2090 2095 2100  
 RELES SVSFP ADITK PPERK EGPVV RLTRR RTTFL TAQPE PRAER ETSSS  
 2105 2110 2115 2120 2125 2130 2135 2140 2145 2150  
 RRRRH PDEPG QFAVA LVVTY RTLGQ LLPEH YDPDH RSLRL PNRPV INTFV  
 2155 2160 2165 2170 2175 2180 2185 2190 2195 2200  
 VSMV YSEGT FLPSL LQRPI LVEFS LLEYE ERSKP VCVFW NISLD TGGTC  
 2205 2210 2215 2220 2225 2230 2235 2240 2245 2250  
 GWSAK GCELL SRNRT HVTQY CSHSA SCAVL MDISR REHGE VLPLK IITYA  
 2255 2260 2265 2270 2275 2280 2285 2290 2295 2300  
 ALSLS LVALL VAFVL LSLVR TLRSN LHSIP QEPFH ALFFS QLIFM VGINQ  
 2305 2310 2315 2320 2325 2330 2335 2340 2345 2350  
 TENPF LCTVV AILLH YVSMG TFAWT LVENL HUYRM LIEVR NIDTG PMAFY  
 2355 2360 2365 2370 2375 2380 2385 2390 2395 2400  
 TM4 HVAGW GIPAI VTGLA VGLDP QGYGN PDFCW LSLQD TLIWS FAGPV GTVII  
 2405 2410 2415 2420 2425 2430 2435 2440 2445 2450  
 INTVI FVLSA KVSQY RRGHY YERKG VVSMK RTAFL LLLLV TATWL LGLLA  
 2455 2460 2465 2470 2475 2480 2485 2490 2495 2500  
 VNSDT LSFHY LFAAF SCLQG IFVLL FYCVA NREVR KHLRA VLAKK KLQLD  
 TM5 TM6

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2505	2510	2515	2520	2525	2530	2535	2540	2545	2550
DSATT	RATLL	TRSLN	CNNVTY	SEGSR	HAPHR	PQOST	ASLDS	TTRDE	GVQKL
2555	2560	2565	2570	2575	2580	2585	2590	2595	2600
SVSSG	PAPGN	HGEPD	ASFIP	RNSKK	ANGPD	SDSDS	ELSLD	EHSSS	YASSH
2605	2610	2615	2620	2625	2630	2635	2640	2645	2650
TSOSE	ODGGE	AEDKW	NPAGG	PAHST	PKADA	LANHV	PAGWP	DESLA	GSDSE
2655	2660	2665	2670	2675	2680	2685	2690	2695	2700
ELDTE	PHLKV	RPRSA	WSYTG	RRRAI	TVATG	PLTRK	VGSWP	SQWPC	LAASP
2705									
RSSGK	AS								

FIG. 10 CONTINUED